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RESULT 2557
ADD81659
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                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                               is useful for predicting the potential of an oligonucleotide to hybridise to a target nucleotide sequence, e.g. RNA or DNA or a sequence that contains chemically modified nucleotides. The method is also useful for predicting the potential of the oligonucleotides to hybridise to a complementary target nucleotide sequence. The method is useful to predict efficient hybridisation oligonucleotides for each of multiple target sequences therefore very large arrays may be constructed and tested with minimum synthesis of oligonucleotides. The present sequence represents a HIV PRT antisense derived probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Predicting potential of oligonucleotides to hybridize to target nucleotide sequence comprises determining and evaluating for each oligonucleotide a parameter predictive of the oligonucleotides ability
Shannon
                                                                                                                                                                                                                          ΛIΗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shannon KW, Wolber PK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (WOLB/)
                   (SHAN/)
(WOLB/)
(DELE/)
(WEBB/)
(KINC/)
                                                                                                                                                 US2003054346-A1
                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                        ADD81659
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hybridize
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-FEB-1998;
                                                                                    10-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (KINC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NAHE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ligonucleotides
                                                                                                                                                                                                   oligonucleotide hybridisation
                                                                                                                                                                                                                       PRT antisense derived probe #588.
                                                                                                                                                                                                                                                                                                                                                                5700
                                                                                                                                                                    immunodeficiency virus
                                                                                                                                                                                          array;
                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
17; Conserv
                             SHANNON K W.
WOLBER P K.
DELENSTARR G
WEBB P G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WOLBER P K.
DELENSTARR G
WEBB P G.
 3
                     KINCAID R H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KINCAID R H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SHANNON
                                                                                                                                                                                                                                                                                                                                           TIGCCTICCTTITCCTCTTC
                                                                                                                                                                                                                                                                                       standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 1 A; 8 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                          minimum
                                                                                                        2001US-00784674
 Wolber PK,
                                                                                                                                                                                                                                            (first entry)
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                                                                                    98US-00021701
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to hybridise to target
                                                                                                                                                                                                                                                                                        DNA,
                                                                                                                                                                                                                                                                                                                                                                                              0.2%;
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                                          a
                                                                                                                                                                                        nybridisation potential; efficient hybridisation; oligonucleotide synthesis; probe.
                                                                                                                                                                                                                                                                                        20
 Delenstarr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Delenstarr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              423pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method of predicting the potential ise to target nucleotide sequences.
                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                               Score 15.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                 5719
                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                                              11 T; 0
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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ဂ္ပ
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                                                                                                                                                                                                                                                                                                                                                                                                .7e+03
 Webb
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                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
g,
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Kincaid
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The method
                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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oligonuclectides to hybridise to target nuclectide sequences. The method is useful for predicting the potential of an oligonuclectide to hybridise to a target nuclectide sequence, e.g. RNA or DNA or a sequence that contains chemically modified nuclectides. The method is also useful for predicting the potential of the oligonuclectides to hybridise to a complementary target nuclectide sequence. The method is useful to predict efficient hybridisation oligonuclectides for each of multiple target sequences therefore very large arrays may be constructed and tested with minimum synthesis of oligonuclectides. The present sequence represents a HIV PRT antisense derived probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Predicting potential of oligonucleotides to hybridize nucleotide sequence comprises determining and evaluationigonucleotide a parameter predictive of the oligonuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hybridize with target.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; SEQ ID NO 732; 423pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and evaluating for each the oligonucleotides ab
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ability to
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밁 8 Query Match Best Local S Matches 17 Sequence 5699 _ Similarity 20 rrreccrrecrrrecrer 5718 BP; 1 Conservative A; 7 0.2%; C; 0 0, G; 12 T; Score 15.2; DB 1; Pred. No. 1.7e+03; 0; Mismatches 3 20 0 Ģ DB 1; 0 Other; u --Length 20; 0, Gaps

0

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RESULT 2558
                                                                       ss; single nucleotide polymorphism; immunosuppressive; antidiabetic; neuroprotective; antirheumatic; antiarthritic; thyromimetic; antiarteriosclerotic; antiinflammatory; dermatological; antipsoriatic; antiasthmatic; diagnosis; autoimmune disease; ESE-3; ESE-2; ESE-1; diabetes; multiple sclerosis; rheumatoid arthritis; lupus; psoriasis; asthma; myasthenia gravis; Sjogren's syndrome; Hashimoto's thyroiditis; Pemphigus vulgaris; atherosclerosis; rheumatoid arthritis; restenosis;
Homo
                                                                                                                                                                                                                                                                                               ESE
                                                                                                                                                                                                                                                                                                                                                   29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                        ADE50858
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE50858
                                                                                                                                                                                                                                                                                               gene
sapiens.
                                                                                                                                                                                                                                                                                               SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA; 20
                                                                                                                                                                                                                                                                                               primer
                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                               #16.
                                                                                                                                                                                                                                                                                                                                                   entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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WO2003034896-A2

01-MAY-2003

15-OCT-2002; 2002WO-US032116

12-OCT-2001; 26-APR-2002; 2001US-0329158P. 2002US-0376139P.

(BETH-) BETH ISRAEL DEACONESS MEDICAL CENT

Libermann 'n Tautu Ó Grall <u>'</u>E ည ×

2003-441218/41

Diagnosing the presence, predisposition or susceptibility autoimmune disease e.g. diabetes or multiple sclerosis, condetecting a polymorphism in the ESE-3, ESE-1 or ESE-1 general sections.

8x55555555555555555555

20

BP; 9 A;

4 C; 4 G;

3 T; 0

U; 0 Other;

DB 1;

polymorphism in the ESE-3, ESE-2 or ESE-1 genes, which is correlated with an alteration in the activity or expression of a polypeptide encoded by these genes. Detection of the polymorphism is indicative of the occurrence, predisposition or susceptibility to autoimmune disease. The method is useful for diagnosing the presence, predisposition to, or susceptibility to an autoimmune disease, e.g. diabetes (e.g. Type I diabetes or Type II diabetes), multiple sclerosis, rheumatoid arthritis, lupus, psoriasis, asthma, myasthenia gravis, Sjogren's syndrome, Hashimoto's thyroiditis, Pemphigus vulgaris, or inflammation (e.g. the atheroselerosis, rheumatoid arthritis, or inflammation associated with restenosis). The method is also useful for preventing or treating any of these diseases. This sequence corresponds to a primer used in the method to detect the single nucleotide polymorphisms in the ESE genes,

predisposition or a susceptibility to the disease, polymorphism in the ESB-3, ESE-2 or ESE-1 genes, \mathbf{w}

The invention relates to the diagnosis of

an autoimmune

γď

detecting a

disease,

Example 2;

SEQ ID NO 72; 96pp; English

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ADE43461
ID ADE8
XX ADE4
XX ADE4
XX ADE4
XX ADE4
XX ADE4
XX Huma
XX Huma
XX AL121
XX AL21
XX AL21
XX AL21
XX ADE4
XX A
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Best Local (
                                                                                                                                                                                                                                   Becker KD, vo
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09-NOV-2001;
04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neurodegenerative disease; uPA; SNCG; Alzheimer's disease; neuroprotective; Chromosome 10; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human SNCG sequencing primer, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE43461 standard; DNA;
                                                                               Determining a predisposition for or the occurrence of neurodegenerative disease, e.g. Alzheimer's disease by detecting in a target nucleic acid the presence or absence of an allelic variant of one or more polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-OCT-2001;
08-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003054143-A2
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                                                                                                                                                                                                                                                                                                                              (NEUR-)
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17; Conser
                                                                                                                                                                                                                                                                                                                              NEUROGENETICS INC.
GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTTTGGCTCATGTGGATTTG 5600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.2%;
nilarity 85.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 ; 2001US-0339525P.
; 2001US-0336929P.
; 2001US-0338010P.
; 2001US-0338363P.
; 2001US-0337052P.
                                                                                                                                                                                                                                              Velicelebi G,
Mullin KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry
                                                                                                                                                                                                                                                                                                                                                                                                             2002US-0368919P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                                                                                                                                                                                                           Elliott KJ,
Sampson AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВÞ
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pred. No. 1.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IDE; KNSL1; LIPA; TNFRSF6;
nootropic; gene therapy;
                                                                                                                                                                                                                                              Wang X, T
Blacker DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.7e+03
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                                                                                                                                                                                                                                                                           Tanzi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                           RE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  °.
                                                                                                                                                                                                                                                                           Bertram
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X000000000000000000X
                                                                             The present invention relates to a method (M1) for determining a CC predisposition for or the occurrence of neurodegenerative disease in a CC subject. The method comprises detecting in a target nucleic acid obtained from the subject the presence or absence of an allelic variant of one or CC more polymorphic regions of one or more genes selected from upA (Urokinase plasminogen activator), SNGG (gamma-synuclein), IDE (insulin-CC lypase), and TNFRSF6 (Tumour Necrosis Factor Receptor-SP6), where the presence of at least one of the allelic variant of one or more CC polymorphic regions is indicative of a predisposition for or the CC cocurrence of neurodegenerative disease. The genes are all located on CC chromosome 10, M1 is useful for determining a predisposition for or the occurrence of, and for treating neurodegenerative disease, particularly Alleimer's disease. The present sequence is a PCR primer, which was used
Sequence 20 BP; 6 A; 1 C; 11 G; 2 T; 0 U; 0 Other;
                                                                Alzheimer's disease. The presen in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                               acid obtained
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RESULT 2560
AAQ29595/c
                                                                                                                                                                                      Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                   Query Match
Best Local :
                                                                                AAQ29595
Pol 67/70 region sequencing primer 90-417.
                        25-MAR-2003
10-MAR-1993
                                                                                                                                                                 3627 GGGGGTGGGAGGAGGTAG 3646
                                                                                  standard; DNA; 21 BP
                                                                                                                                                                                        Conservative
                       (revised)
(first entry)
                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                   Score 15.2;
Pred. No. 1
                                                                                                                                           20
                                                                                                                                                                                                  .7e+03
                                                                                                                                                                                                                DB 1;
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                                                                                                                                                                                          Indels
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                                                                                                                                                                                          Gaps
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밁 Ś

Synthetic Amplify; HIV; pol; resistance; azidothymidine; AZT; 3SR; probe; self-sustained sequence replication; mutation; inosine; ss.

WO9216180-A2

12-MAR-1992; 92WO-US002037

13-MAR-1991; 03-SEP-1991; 91US-00668549 91US-00754146

SISKA UNIV (CALIFORNIA.

Gingeras TR, Barringer KJ, Richman DD, Prodanovich 'n, Davie GR;

WPI; 1992-348902/42

Assay for detecting genotype of AZT resistance - utilises probes which anneal to amplified region of HIV-1 gene. series o m

Disclosure; Page 17; 57pp; English.

The sequences given in AAQ29591-601 are sequencing primers which were used in the method of the invention to determine the sequence of regions of the HIV pol gene which are involved with resistance to azidothymidine (AZT). Resistance to AZT is caused by the accumulation of four mutations grouped within two regions of the HIV pol gene. The primers AAQ29591-96 are used to determine the sequence around the mutations at amino acids positions 67 and 70, and primers AAQ29597-601 are used to determine the sequence around the model to determine the sequence around the mutations at a mino acids positions 67 and 70, and primers AAQ29597-601 are used to determine the

Example 2;

Page

267;

848pp; English

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RESULT 2561
AAQ79211/c
ID AAQ7921
XX AAQ7921
XX GANGR
DT 17-JUL-
XX GUANOSI
KW GUANOSI
KW GUANOSI
KW HEPSTLIT
KW CIAIM
KW HEPSTLIT
KW HEPS
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Best Local
                                           The oligonuclectides (See AAQ79201-52) can be used to treat viral infections. The oligonuclectides inhibit viral replication by forming guanosine tetrads which form a stabilised 3D structure. Preferred oligonuclectides contain at least 2 runs of at least 2 guanosine bases and may be capped at the 3' terminus with a modifier selected from polyamine, poly-lysine, cholesterol and propanolamine. They may also have a modified phosphodiester linkage or be modified to contain a phosphorothicate linkage. They are used to treat infections with viruses such as herpes simplex virus, human papilloma virus, Epstein-Barr virus, HTV, adenovirus, respiratory syncitial virus, hepatitis B virus or human cytomegalovirus. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interest
(Updated
                                                                                                                                                                                                                                                                                                                                                                                         Oligo-nucleotide(s) rich in guanosine which to treat viral infections, e.g. herpes-virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TRIP-)
(BAYU )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-APR-1993;
28-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                              Claim 41; Page 49; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-357890/44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-APR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      infection; herpes simplex virus; human papilloma
Epstein-Barr virus; HIV, adenovirus; respiratory
hepatitis B virus; human cytomegalovirus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Guanosine rich oligonucleotide used to treat viral infection
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17-JUL-1995
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BAYLOR C
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COLLEGE MEDICINE.
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RESULT 2563 AAT51629/c

AAT51629

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Best Local
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                                                                                                       AAT16454-T16469 are PCR primers used to amplify human p53 genomic DNA. The primers are used to demonstrate a method of sequence-based diagnosis of a human neoplastic tissue, blood or other body fluid. The method determines the presence, nature and location of any mutations and their influence on the biological function of the p53 protein (or other cancerrelated protein), and hence the properties of the neoplasia (e.g. metastatic potential). The method allows a reliable, accurate diagnosis of neoplasia, allowing clinicians to form a prognosis on the development of the neoplasia and guidance for its treatment
                                                                                       Sequence 21
                                                                                                                                                                                                                                             Sequence-based diagnosis of a human neoplastic tissue, blood fluid - useful for the diagnosis or prognosis of neoplasia.
                                                                                                                                                                                                                                                                                                                                                                                29-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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16-NOV-1994;
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94SE-00003953
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Pred. No. 1.8e+03;
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RESULT 2564
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUL-1995;
23-OCT-1995;
19-MAR-1996;
25-MAR-1996;
17-APR-1996;
23-APR-1996;
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  promoter; template probe; signal amplifier; hybridisation assay; detect; functional domain; DNA-dependent RNA polymerase; quantifying; analyte; ligand receptor; amplification; hepatitis B virus; Neisseria gonorrhoeae
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                                                                                                                                          RESULT 2565
AAT95441/c
                                                                                                                                                                                                                                                                                                                                         CC A novel DNA construct (referred to as a "template probe") for use as a CC signal amplifier in hybridisation assays to detect a target comprises 3 CC functional domains (A, B and C) orientated A-B-C or B-C-A. (A) is single-CC stranded and is designed to hybridise to complementary target sequence CC promoter. (C) is single-or domain B. It consists of a nucleotide CC for the promoter activity of domain B. It consists of a nucleotide CC sequence not found in the template. The DNA construct is used in a method CC for detecting and quantifying an oligonucleotide analyte or a ligand CC receptor by amplification of a biological signal in a nucleic acid CC nucleic acid segments characteristic of hepatitis B virus, Neisseria CC nucleic acid segments characteristic of hepatitis B virus, Neisseria CC microtioneae, bacterial beta-lactamase TEM-1 gene, Chlamydia, bacterial CC the pilin gene DNA of Neisseria gonorrhoeae was performed using a CC microtitre dish assay procedure and the T7 RNA polymerase. The present CC con 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 17
                                                     25-MAR-2003
10-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bacterial beta-lactamase TEM-1 gene; Chlamydia; HIV; hepatitis C virus; bacterial tet M determinant; consensus; T7 promoter; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA construct for use as signal amplifier in hybridisation assays containing DNA-dependent RNA polymerase promoter and template sequ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Urdea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAR-1994;
                          Primer for breast cancer susceptibility gene BRCA2 exon 11-7.
                                                                                                  AAT95441;
                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-280266/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JAN-1990;
10-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-MAY-1997.
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                                                                                                                              AAT95441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHIR ) CHIRON
                                                                                                                                                                                                                     3609 TTCTTTGGGGAATGGGGTGG 3628
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                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Col 25; 45pp; English
                                                                                                                             standard; DNA; 21
                                                                                                                                                                                                    TTCTTTGGAGAAAGTGGTGG 1
                                                                                                                                                                                                                                                                                                                  BP; 7 A; 8 C; 2 G; 4 T; 0 U;
                                                                                                                                                                                                                                                             Conservative
                                                       (revised)
(first ent
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91US-00639560
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/note= "N4-(6-aminocaproyl-2-aminoethyl) derivative of 5-
methyl cytidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                        entry)
                                                                                                                                                                                                                                                                       0.2%;
                                                                                                                              ВP
                                                                                                                                                                                                                                                             ;
                                                                                                                                                                                                                                                                         Score 15.2; DB 1 Pred. No. 1.8e+03
                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                    0 Other;
                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                             Gaps
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Human; breast cancer; susceptibility; gene; BRCA2; diagnosis; screening;

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RESULT 2566
AAT80147/c
ID AAT80147;
XX AAT80147;
AC AAT80147;
XX Immunoglob
XX Immunoglob
XX CTLA4; Igc
KW diabetes in
KW diabetes in
KW transplani
KW transplani
KW transplani
KW immunosupp
XX MO9728267;
XX WO9728267;
XX O7-AUG-19;
XX O3-FEB-19;
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20-DEC-1995;
21-DEC-1995;
11-JAN-1996;
29-APR-1996;
                                                                                              myasthenia gravis; systematransplant rejection; graftransplant rejection; human;
                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is a primer for the human breast cancer susceptibility gene BRCA2, which can be used to diagnose breast cancer and screen for compounds to treat breast cancer BRCA2 can also be used in gene therapy to restore wild type BRCA2 gene function to a cell, which has lost its or has altered (i.e. by virtue of a mutation in BRCA2) BRCA
                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                           gene function.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-341680/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MYRI-)
(UYPE-)
(HSCR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-1997
 03-FEB-1997;
                       07-AUG-1997.
                                             WO9728267-A1
                                                                                                                                    diabetes mellitus;
                                                                                                                                               CTLA4; IgG1;
                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tavtigian SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9722689-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment; gene therapy; PCR primer; exon 11-7; ss.
                                                                                                                                                                    Immunoglobulin signal sequence reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENDO-)
                                                                                                                                                                                                                                                                                                                                                    Loca1
                                                                                                                                                                                                                                                                                                                    4947
                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                          17; Conserv
                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer
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HSC RES & DEV LP.
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                                                                                                                                                                                                                                                                                                           TTACTTTTTTCCTGCTGGCT 4966
                                                                                                                                                                                                                                                                                                                                                                                      21 BP;
                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                              TAACTTTTTTCCCGCTAGCT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 60; 189pp; English.
                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                          immunoglobulin; antibody; autoimmune disease;
litus; rheumatoid arthritis; multiple sclerosis;
ravis; systemic lupus erythematosus; thyroiditis;
ejection; graft versus_host disease; allergy; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer
                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and
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95US-00575359.
95US-00576559.
96US-00585391.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kamb
 97WO-US001698.
                                                                                                                                                                                                                                                                                                                                                                                                           (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-00639501
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                                                                                                                                                                                                                                                                                                                                                                                      9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            er susceptibility gene BRCA2 screening for compounds to
                                                                                                                                                                                                                                       DNA;
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                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                    3 C; 6 G;
                                                                                                                                                                                                                                       21
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                                                                                                  primer;
                                                                                                                                                                                                                                                                                                                                          0
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                    3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ŗ
                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                   PCR;
                                                                                                                                                                                                                                                                                                                                                     15.2;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Couch
                                                                                                 signal peptide;
                                                                                                                                                                    PCR primer.
                                                                                                                                                                                                                                                                                                                                                    .8e+03
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treat breast c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rommens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer.
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                                                                                                                                                                                                                                                                                                                                         Gaps
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BRCA2
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RESULT 2567
AAX79218/c
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A forward primer (AAT80146) and a reverse primer (AAT80147) were used for the PCR amplification of the immunoglobulin (Ig) signal sequence from template plasmid pSP72IGG1. The reverse primer corresponds to the C-terminal of the natural Ig signal peptide. The 208 bp PCR product contains the entire Ig signal sequence. Novel CTLA4-antibody fusions have the heavy and light chain variable domains of an antibody molecule replaced with the extracellular domain of CTLA4. The resulting antibody-like protein binds to B7-1, B7-2 and CTLA4 ligands with high affinity. Novel CTLA4-Ig constructs (see AAT80131-83) encode fusion proteins (see AAW26206-08) useful in claimed methods for suppressing an immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gray
                                                                                                                                                                               Key
modified_base
                                                                                                                                  misc_structure
                                                                                                                                                                                                                                                 herpes simplex virus; HSV; human papilloma virus; Epstein Barr Virus; HPV; EBV; HIV; human immunodeficiency virus; ademovirus; RSV; HBV; HCM respiratory syncytial virus; hepatitis B virus; human cytomegalovirus; human.T-cell leukaemia virus; HTLV; ss.
                                                                                                                                                                                                                                                                                                                                                                                                    AAX79218 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New CTLA4-modified immunoglobulin fusion proteins - treating auto:immune diseases and allergies, or for
                                              06-AUG-1998.
                                                                       W09833807-A1
                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                  Column;
                                                                                                                                                                                                                                                                                                                          Oligonucleotide #11 forms an intramolecular stacked tetrad structure.
                                                                                                                                                                                                                                                                                                                                                     31-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                            AAX79218;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 46; 105pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transplantation rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in a subject
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6020
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TCTCCACAGGTGTCCACTCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTCCACACCTGTCCACTCC 6039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 5
                                                                                                                                                                                                                                                                                                  stacked tetrad; inhibition; replication; pathophysiological;
                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96US-00595590
                                                                                                                                                                                               Location/Qualifiers
                                                                                                 columnar
                                                                                                                                              /*tag= a
/note= "optionally contains
internucleotide linkages"
                                                                                                          /*tag= b
/note= "forms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; 3 C; 10 G; 3 T;
                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                                                 box structure"
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Pred. No. 1.
                                                                                                            intramolecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
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                                                                                                                                                         phosphodiester
                                                                                                            stacked tetrad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ω,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rennert PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
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                                                                                                                                                                                                                                                                           HCMV;
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04-FEB-1997; 03-FEB-1998;

97US-0037374P 98WO-US001974

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AAZ26573/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequences AAX79210-X79275 represent oligonucleotides (ON) which are able to form a columnar box or "stacked tetrad" structure by intramolecular internucleotide binding. The ONs are used to inhibit the replication of viruses. They are able to suppress virus production for prolonged periods after an initial short treatment regimen. They can be used for treating pathophysiological states caused by viruses such as herpes simplex virus, human papilloma virus, Epstein Barr Virus, HIV, adenovirus, respiratory syncytial virus, hepatitis B virus, human cytomegalovirus and HTLV I and
                                                                                                                                                                                                                                                                                          Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; pervention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New guanosine-rich tetrad forming oligonucleotide(s) - used for inhibiting virus replication for treating e.g. herpes simplex, papilloma HIV, adenovirus or hepatitis B virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
                         prevention dysplastic
                                                                                                                                                                                                               24-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                     Human polymorphic region 762
                                                                                                                                                                                                                                                                                                                                                                                                                  30-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ26573
                                                                               WPI; 1998-521232/44.
                                                                                                                                                           20-MAR-1997;
                                                                                                                                                                                     19-MAR-1998;
                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ARON-) ARONEX PHARM
                                                                                                       Housman D,
                                                                                                                                 (VARI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2998
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                                                                                                                                 VARIAGENICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
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                                        and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 137;
                                                                                                     Ledley FD,
                                                   target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                      rget genes
treatment
                                                                                                                                                          97US-0041057P
                                                                                                                                                                                      98WO-US005419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 0 C; 17 G; 4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2
85.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hogan ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 BP
                                                                                                        Stanton
                                     for allele-specific drugs - used for diagnosis, of, e.g. cancers, atherosclerotic plaque,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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Pred. No. 1.8e
O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wallace
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3; Indels
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Disclosure; Fig 7;

605pp; English

lesions, endometriosis

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graft versus host disease

This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is acon a gene vital for cell growth or viability, and where the gene is

active

Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.

WPI; 1998-521232/44.

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Ledley

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Stanton

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(VARI-)

VARIAGENICS

Disclosure;

Fig 7; 605pp; English.

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This invention describes a novel method for identifying an inhibitor CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient a first allele greeinfic inhibitor (ASI) targeted to an allele of a first essential gene cC present in cells of the precancerous condition, where the normal somatic CC cells of the patient are heterozygous for the first gene, the inhibitor CC calls of the patient are heterozygous for the first gene, the inhibitor CC calls of the patient are heterozygous for the first gene, the inhibitor CC cells of the gatient are heterozygous for the first gene, the inhibitor CC cancers in a population and targets only one allelic form present in the CC cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic CC lesions, benign tumours, endometriosis, polycystic kidney disease, and CC graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent contacts.
Sequence 21
      BP;
      14 A; 1 C; 3 G; 3 T; 0 U; 0 Other;
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RESULT 2569
AAZ26268
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Matches
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20-MAR-1997;
                                     19-MAR-1998;
                                                                          24-SEP-1998.
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17; Conserv
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97US-0041057P
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85.0%;
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Pred. No. 1.8e+03;
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RESULT 2570
AAZ26511
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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is action a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele
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                                                                                                                                                                                                            Disclosure; Fig
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or graft
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used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene

This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is ac on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is

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                                                                                                                                                                                                Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
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RESULT 2572

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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is considered to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) traggeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic form present in the normal somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic collesions, benign tumours, endometriosis, polycystic kidney disease, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ26485 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-521232/44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig .7; 605pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5744
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VARIAGENICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ledley FD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0041057P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-US005419
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Pred. No. 1.8e+03;
D; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
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                                                                                 CC This sequence represents a PCR primer for a barley chromosome 7 CC microsatellite marker, and can be used in the method of the invention. CC The method is for screening cereal for fermentability, comprising careal for screening cereal for fermentability, comprising careal for screening chromosome of a gene/gene CC complex affecting fermentability at a locus close to the centromere on the homologous Triticeae group 5 chromosome (barley chromosome 7) is/are CC present. The invention also relates to a method for screening cereal for CC comprising analysing barley genomic DNA to determine which allele(s) of the locus, designated eph on the short arm of homologous Triticeae group 1 chromosome (barley chromosome 5) is/are present. The methods and CC primers are useful for identifying microsatellites Bmac96 and Bmac213, which are useful for determining fermentability and/or epi-heterodendrin production in cereals, especially barley. Current methods for determining fermentability are difficult to apply within barley breeding programs.

CC Prior art methods using molecular markers have difficulty in detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 25; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thomas WTB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SCCR-) SCOTTISH CROP RES INST
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17; Conserv
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                                                            methods using molecular markers have difficulty allelic variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cereals for fermentability, especially useful in barley.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             microsatellite marker; barley; chromosome 7 marker; cereal; ity; group 5 chromosome; ethyl carbamate production; Bmac213 tation; Triticeae; Bmac96; epi-heterodendrin production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer for microsatellite marker clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rom bone marrow transplants. AAZ25 sites described in the method of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Powell W,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Waugh R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .8e+03;
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Query Match Best Local Similarity

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Score Pred.

15.2; DB 1; No. 1.8e+03;

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                                              CC CNS glial-restricted precursor (GRP) cells which can self-renew in the CC AFCI culture medium and can differentiate to CNS glial cells but not to CC CNS enuronal cells. The stem cells can be used to generate a population CC computing cell proliferation and neuronal differentiation. The medium CC promoting cell proliferation and neuronal differentiation. The medium CC comprises laminin-coated plates and NEP medium lacking chick embryo CC extract. The stem cells can also produce neural crest stem cells by CC inducing the cells to differentiate in vitro. The inducing step comprises CC replating the cells on a laminin-coated substrate and preferably CC withdrawing a mitogen (preferably fibroblast growth factor; FGF) and CC chick embryo extract. Inducing can also comprise adding a dorsalizing CC agent to the cells, preferably a bone morphogenetic protein (BMP) such as CC BMP-2, -4 or -7. The stem cells can be used to produce cells of the CC vitro to neural crest stem cells can be used to produce cells of the CC vitro to neural crest stem cells, and inducing these cells to differentiate in CC comprises adding the stem cells to an example from the present invention to amplify different FGF and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated, pure population of mammalian neurospithelial stem cells, which are capable of self-renewal in adherent feeder-cell-independent (APCI) culture medium and differentiation to central nervous system (CNS) neuronal or glial cells and to neuronal crest stem cells. Also described is an isolated population of mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalian neuroepithelial stem cells and glial restricted precursor - self renew and differentiate into central nervous system cells, used generating various types of cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neuroepithelial stem cell; lineage restricted intermediate precursor; oligodendrocyte; astrocyte; self-renewal; neuron; differentiation; neural creat cell; fibroblast growth factor; FGF; FGFF; receptor; CNS; central nervous system; glial cell; PCR primer; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-070093/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rao MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAY-1997;
06-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-MAR-1999
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                                  FGFR genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 59; 78pp; English.
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98US-00073881.
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Query Match

Sequence

21 BP; 6

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4 C; 6

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04-SEP-2000 AAA46283;

(first entry)

AAA46283

AAA46283 standard; DNA; 21

ВP

RESULT 2576

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Length

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AAZ95022
                                                   Matches
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                                                                            Query Match
                                                                                                                          The present sequence is that of the forward primer used in the real-time quantitative PCR amplification of cancer specific gene Pro111 (see AAZ95002 and AAZ95003). Overexpression of Pro111 was found in 5 of 16 primary prostate cancer samples examined, indicative of it being a diagnostic marker for prostate cancer. The invention provides ESTs and full-length contigs for CSGs (see AAZ94998-Z95017). The CSGs, polypeptides encoded by them, and antibodies that specifically bind CSG are used in claimed methods for detecting, diagnosing, monitoring, staging, imaging and treating prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prostate cancer; cancer specific gene; CSG; expressed sequence tag; Ediagnosis; monitoring; staging; imaging; therapy; metastasis; marker; human; Proll1; PCR primer; ss.
                                                                                                      Sequence 21 BP; 5 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         measuring changes in cancer specific gene levels.
                                                                                                                                                                                                                                                                                                       Diagnosing, staging and monitoring the presence and metastases prostate cancer especially useful for treating prostate cancer
                                                                                                                                                                                                                                                                                                                                               WPI; 2000-339531/29.
                                                                                                                                                                                                                                                                                                                                                                          Salceda
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prostate cancer diagnostic marker Proll1 forward PCR primer.
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                                                                                                                                                                                                                                                                  Example
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                                                                                                                                                                                                                                                                                                                                                                                                    (DIAD-) DIADEXUS LLC.
                                                               Local
                        5181 CTGCATGTTCTCCACTTGGA 5200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3683 GCCAGAAAGCCAGCTATTTT 3702
                                                 cal Similarity
17; Conserv
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                                                                                                                                                                                                                                                                  2; Page 23; 74pp; English.
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CTGCAAGTTCTCCACATTGA 21
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                                                              0.2%;
85.0%;
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                                                 0; Mismatches
                                                               Score 15.2;
Pred. No. 1
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                                                                                                                                                                                                                                         real-time
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RESULT 2577
AAZ7528)/c
ID AAZ7528
XX
AC AAZ7528
XX
DT 10-SEP-
XX
DE Human b
XX
KW Human 9
KW Human 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC interphotoreceptor matrix (IPM) proteoglycan, designated IPM200. The protein is an IPM component (IPMC). Two subfamilies of IPMCs, IPM150 and CC protein is an IPM component (IPMC). Two subfamilies of IPMCs, IPM150 and CC IPM200, exist. The human IPM150 gene is located on chromosome 6q13-q15, component a patients own production of the protein or to rectify CC to supplement a patients own production of the protein or to rectify CC alterations in their nucleic acids that result in expression of an CC inactive protein. The IPM nucleic acids may be used in this way to treat CC ocular diseases such as retinal detachment, chorioretinal degeneration, cone CC retinal degeneration, age related macular degeneration, cone CC degeneration, mucopolysaccharidosis, rod-cone dystrophy and cone-rod CC degeneration, mucopolysaccharidosis, rod-cone dystrophy and cone-rod CC dystrophy. The nucleic acids and proteins may also be used to treat ocular diseases. The nucleic acids and proteins may also be used as diagnostic resgents to detect the presence of IPM nucleic acids and proteins from their receiptor and contact of the proteins may also be condition.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interphotoreceptor matrix; IPM; proteoglycan; IPM150; II chromosome 6q13-q15; ocular disease; retinal detachment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acids encoding interphotoreceptor matrix proteoglycans useful preventing, diagnosing and treating ocular disorders such as retinal detachment and chorioretinal degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-365616/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer for interphotoreceptor matrix proteoglycan IPM200 cDNA.
Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 43; Page 120; 183pp; English
                                                                                              Human biallelic marker downstream amplification primer SEQ ID NO:9636.
                                                                                                                                                                                                                                                                                        AAZ75280 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP;
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Pred. No. 1.8e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 21;
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AAZ72608
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-APR-1998;
;23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-OCT-1999.
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                                                                                                                          Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel biallelic markers used to construct a high density disequilibrium
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                                                                                                                                                                                                                                                               Human biallelic marker upstream amplification primer SEQ ID NO:6964.
                                                                                                                                                                                                                                                                                                                                                                           AAZ72608;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21
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WO9954500-A2
                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                   10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ72608 standard; DNA;
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98US-0109732P
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AAZ76810
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention, which contain a polymorphic base at position 24 of their nucleotide sequences. ARZ65579 to ARZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not accually given a sequence in the Sequence Listing from the present invention.
                                                                                                                                                                                                 Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; identification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
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                                                                                                                                                                                                                                                                        Human biallelic marker downstream amplification primer SEQ ID NO:11166.
                                                                                                                                                                                                                                                                                                                                                                AAZ76810
                           21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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23-NOV-1998;
                                                                     21-APR-1999;
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                                                                                                                              WO9954500-A2
                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    map of the human genome.
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(GEST ) GENSET
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98US-0109732P.
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98US-0109732P.
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Pred. No. 1.
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AAC60966/c
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Best Local
                                                                                                                                                                                                                                                                                                                           snort tandem repeat; primer; STR; susceptibility; HIV; detection; polymorphism; interleukin 10 promoter; IL-10 chromosome position 6p21.3; tumour necrosis factor have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not accually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cohen D, Blumenfeld
WPI; 2000-013267/01
                                    Predicting susceptibility to HIV infection or progression useful for selection of therapeutic treatment for persons infected with HIV viracomprises detecting polymorphism in human interleukin-10 promoter.
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                                                                                                                                                                                                                                                                                                                                                                                        Tumour necrosis factor beta short tandem repeat primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC60966;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the in
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                                                                                                                                                       (USSH ) US DEPT HEALTH & HUMAN SERVICES.
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                                                                                                                                                                                                                                                                                                    Homo sapiens.
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                 nism; interleukin 10 promoter; IL-10; 6p21.3; tumour necrosis factor beta; ss
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Pred. No. 1.8e+03
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Example 1; Page 12; 40pp; English.

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RESULT 2581
AAC80269/c
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Best Local
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The present invention relates to a method for the locus-specific, separate amplification of exon 2, exon 3, and/or exon 4 of human leukocyte antigen (HLA)-A, HLA-B, or HLA-C alleles. The method is useful for subtyping or typing of HLA class 1 alleles. The present sequence is an amplification primer used in the method
                                                                                                                                                                                                                        Locus-specific, separate amplification of exon 2, exon 3, and/or exon 4 of human leukocyte antigen (HLA)-A, HLA-B, or HLA-C alleles using defined primer sets, useful for subtyping or typing of HLA Class I alleles.
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                                                                                                                                                                       Claim 4; Page
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11-JUN-1999;
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17; Conserv
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99US-0138614P.
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Query Match Best Local

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Conservative

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Score 15.2; D Pred. No. 1.8e 0; Mismatches

1.8e+03

Length

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RESULT 2582
AAF95320/c
ID AAF9532
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Best Local S
Matches 17
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26-JUL-2000;
16-AUG-2000;
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                                                                                                                                             The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart
                                                                                                                                                                                                                                                                                                                                                          Nucleic acids comprising single nucleotide polymorphisms, useful applications such as forensics, paternity testing, medicine, gene analysis and phenotype correlations to diseases such as diabetes
                                     disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human gene single nucleotide polymorphism #81.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lander ES,
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2000US-0225724P.
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Pred. No. 1.8e+03;
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RESULT 2583
AAF95430/c
ID AAF9543
XX AAF9543
XX AAF9543
XX AAF9543
XX AAF9543
XX Human 9
XX W020011
XX W020011
XX W020011
XX Homo 86
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XX Human 9
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Matches 17
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26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                               genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids comprising single nucleotide polymorphisms, us applications such as forensics, paternity testing, medicine, analysis and phenotype correlations to diseases such as diab
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                                                                                                                                                                                                                        Sequence
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                                                                                                            l Similarity
17; Conserv
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                                                      AACATCAGCAGCAGCAGCAG 7427
                                                                                                                                                                                                                  21 BP; 0 A; 7 C; 4 G; 10 T; 0 U; 0 Other;
  AACAAGAGCAGAAGCAGCAG
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2000US-0220947P.
2000US-0225724P.
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Pred. No. 1.8e
0; Mismatches
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                                                                                                                                   .8e+03
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RESULT, 2584
AAF96888/c
ID AAF9688
XX AAF9688
XX AAF9688
XX Human g
XX Human g
XX Human g
XX Myolymor
KW Human;
KW polymor
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XX Homo sa
XX Homo s
                                                                                              Query Match
Best Local Similarity
Matches 17; Conserv
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26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosolerosis, myocardial infarction, coronary heart diagnosis of atherosolerosis, myocardial infarction, coronary heart
                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                             pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in foremsics, paternity testing, genetic analysis and phenoty correlations to diseases. The present sequence is an example of one the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (WILL-)
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                                                                                                                                                                                                                                                                                                                                                                    disease, stroke, peripheral vascular diseases, venous thromboembolism and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Page 159; 242pp; English.
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                                               64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WHITEHEAD INST BIOMEDICAL MILLENNIUM PHARM INC.
                              GGCTGCGGGGGGGGCGCGC
                                                                                                                                                                                                             21
                                                                                                                                                                                                             вP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gargill M,
                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0153357P.
2000US-0220947P.
2000US-0225724P.
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                                                                                                                       0.2%;
                                                                                                                                                                                                           13 C; 7
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                                                                                                    0,
                                                                                                                                                                                                           G; 1
                                                                                                                       Score 15.2; DB 1;
Pred. No. 1.8e+03;
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                                                                                                    Mismatches
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                                                                                                                                                                                                           T; 0 U; 0 Other;
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                                                                                                                                                      DB 1;
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                                                                                                                                                                                                                                                                                      phenotype
of one of
                                                                                                    Gaps
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RESULT 2585 AAH22257 ID _AAH2225

, AAH22257 standard; DNA;

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AAH22257;

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CC in angiogenesis (1), and the polypeptides that encode them. (1) have angiogenesis. The nucleic acids and polypeptides may be used in the cardiovascular activity, and can be used in the modulation of angiogenesis. The nucleic acids and polypeptides may be used in the cardiovascular diagnosis and treatment of diseases associated with the production of antibodies against them and in assays to identify and the production of antibodies against them and in assays to identify and antagoniste may also be used as antipodies and activity. The antibodies and activity and contragoniste may also be used to down regulate expression and activity and contragoniste may also be used as diagnostic modulate angiogenesis. The antibodies may also be used as diagnostic constructs for detecting the presence of the polypeptides in samples. The production poliferative tumours, considered with neovascular tumours, proliferative tumours, considered with neovascularisation, peripheral vascular disease. The production, inflammatory vasculitides, Reynaud's disease and considered with neovascularisation, peripheral vascular disease, in phenomenon, aneurysms, arterial restenosis, thrombophlebitis, conditions, heart failure such as congestive heart failure, age-related conditions, heart failure such as congestive heart failure, age-related conditions, heart failure such as congestive heart failure, age-related conditions, chronic heart failure such as congestive heart failure age-related failure, age-related in the exemplification of the present failure such as congestive heart failure, age-related failure failure 
                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1999;
13-APR-2000;
31-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vascular tumour; proliferative tumour; proliferative vitreoretinopathy; rheumatoid arthritis; Crohn's disease; atherosclerosis; endometriosis; neovascularisation; restenneis humania.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; differentially expressed gene; angiogenesis; diagnosis; angiogenic disorder; wound healing; cancer; cardiovascular; ps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Differentially expressed genes involved in angiogenesis, useful treating e.g. vascular tumors, atherosclerosis and/or restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mehraban F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          myocardial infarction; chronic heart condition; osteoporosis; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Placental growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes differentially expressed genes
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                                                                                 Sequence
Local Similarity
les 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 147; 182pp; English
                                                                                 B₽;
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2000US-0196802P
2000US-00703350
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                     Score 15.2;
Pred. No. 1.
                                                                                 6 T;
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                                                                                 0 U;
                     1.8e+03
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                                          DB 1;
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                                      Length 21;
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Matches

Conservative

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RESULT 2587 AAH62143/c ID AAH62143

standard; DNA;

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RESULT 2586
AAH62597/c
ID AAH6259
XX AAH6259
XX AAH6259
XX Single
KW Single

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                                                                                                             Matches
                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                       DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart adiseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; human; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHRNA7 polymorphism containing DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid segments of the human genome, particularly from including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
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                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                      applications such as phenotype correlation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 69; 80pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-NOV-1999;
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21
                                                                                                             1 Similarity
17; Conserv
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                                                                                                                                                                                                                        BP; 1 A;
                                                                                                             Conservative
                                                                                                                                                                                                                                                                            genetic analysis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                           6 C; B G;
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                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                           forensics, paternity testing
                                                                                                                                                                    Length
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RESULT 2588
AAH91825
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                                                                                                                                                                                                                                                                   Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                        DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nuclei acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in
Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
                                                                                      09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
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                                                                                                                                         AAH91825 standard;
                                                                                                                                                                                                                                                                                                                                                                 applications
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 32; 80pp;
                                                                                                                                                                                                                                                                                                                       21 BP; 4 A; 4 C; 6 G; 7 T; 0 U; 0 Other;
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                                                          bowel disease associated
                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                           0.2%;
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Pred. No. 1.8e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                               correlation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             forensic
                                                                                                                                                                                                                                                                             8e+03
                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             science;
                                                        polymorphic site
                                                                                                                                                                                                                                                                                                                                                                 forensics,
                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA fragment #44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation;
                                                                                                                                                                                                                                                                                                                                                               paternity testing
                                                                                                                                                                                                                                                                 <u>.</u>
                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to determine the presence of genetic polymorphisms associated with inflammatory he disease and correlating their occurrence with disease states. They mused in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                           Synthetic
                                                                                                            Sequencing primer; definitive ectoderm equivalent cell; DEE cell; cell preparation; early primitive ectoderm-like cell; EPU cell; hereapy; gene therapy; neuroectoderm cell; organ transplant; neurodegenerative disease; Parkinson's disease; Alzheimer's disease
                                                                                                                                                                                                                                        AAF87033 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 1 A; 3 C; 0 G; 16 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphic site described in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Testing for the presence of polymorphisms associated with bowel disease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-367874/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Daly M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ELLI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           misc_feature
                                                       WO200151610-A1
                                                                                                  stroke;
                                                                                                                                                                      Anchored 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-DEC-1999;
10-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                          4463
                                                                                                                                                                                                                                                                                                                                                1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                   _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WHITEHEAD INST BIOMEDICAL ELLIPSIS BIOTHERAPEUTICS C
                                                                                                  spinal cord injury;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hudson TJ,
                                                                                                                                                                                                                                                                                                                         CTTTTTTTTTTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 76; 463pp; English.
                                                                                                                                                                                                                                                                                                   CTTCTTTATCNTTTTTTTTTT
                                                                                                                                                                     oligo dT12 primer.
                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0196046P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000WO-US033632
                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0170257P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/note= "SNP, optionally T or A at
                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lander ES,
                                                                                                                                                                                                                                         21
                                                                                                  therapy; ss.
                                                                                                                                                                                                                                                                                                                                                 <u>.</u>
                                                                                                                                                                                                                                                                                                                                                         Score 15.2; DB 1;
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                        4483
                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rioux J,
                                                                                                                                                                                                                                                                                                                                                                                                                   exemplification of
                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Siminovitch
                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       this
                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammatory
                                                                                                                                                                                                                                                                                                                                                <u>.</u>
                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                     bowel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detect
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12-JAN-2001; 2001WO-AU000029

19-JUL-2001.

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This sequence represents a sequencing primer used within the scope of the cinvention. The invention relates to a method for preparing definitive ectoderm equivalent (DEE) cells in vitro comprising providing: (a) early CC primitive ectoderm-like (EPL) cells; and (b) a conditioned medium or CC extract exhibiting neural inducing properties and culturing the EPL cells for a time to permit controlled differentiation to DEE cells. The DEE CC cells, or their differentiated or partially differentiated progeny are useful in human cell therapy or transgenic animal production and for use in human or animal gene therapy. The method is useful for preparing DEE Cc cells in vitro. It can also be used for selectively producing neuroectoderm cells or surface ectoderm cells from DEE cells. The method CC useful for preparing tissues or organ for transplant. The cells are useful for preparing tissues or organ for transplant. The cells are creating and curing neurodegenerative diseases such as such as stroke and spinal cord injury by replacing or assisting the treation of normal disease tissues. They are also useful for the cells as source for reprogramming and for use in pharmaceutical or transplant.
RESULT 2590
ABK51835/c
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Best Local
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20-APR-2000; 2000AU-00007045.
27-APR-2000; 2000AU-00007143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Preparing (M1) definitive ectoderm equivalent (DEE) cells in vitro treatment of Parkinson's and Alzheimers comprises culturing early primitive ectoderm-like cells in conditioned medium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               toxicological screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 37; 116pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-432907/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Long CLO,
              07-SEP-2000; 2000JP-00272228
                                              07-SEP-2000; 2000JP-00272228
                                                                                                                                                                                          glucuronic
                                                                                                                                                                                                         Human; enzyme
                                                                                                                                                                                                                                                                        30-JUL-2002
                                                                                                                                                                                                                                                                                                                                      ABK51835 standard;
                                                                                                             JP2002085066-A
                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                         DNA probe #1
                                                                                                                                                                                                                                                                                                        ABK51835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    4458 ATGGACTTTTTTTTTTTT 4477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BRESAGEN
LONG C L
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                                                                                                                                                                                                                                                                                                                                                                                                                    ATGAATTCTTTTTTTTTT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rathjen PD,
                                                                                                                                                                                        /me classification; enzyme quantitative determination;
acid conjugation; UDP-glucuronosyltransferase; UGT8; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 4 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                         for human UGT8 gene
                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.TD.
                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                                                        21 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
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The present invention relates to a method for classification and quantitative determination of enzymes participating in glucuronic acid conjugation. The method involves the use of oligonucleotide probes hybridising to regions of the human UD-glucuronosyltransferase (UGT) genes (e.g. UGT1, UGT1A7, UGT1A9, UGT1A10, UGT2A1, UGT2B7, UGT2B10, UGT2B11, UGT2B15, UGT2B17, UGT8), and the DDOST gene. The method and probes are useful for the genetic determination of enzymes participating in glucuronic acid conjugation with catalysed UGT. The method is both rapid and accurate. ABK51813-ABK51836 represent oligonucleotide probes useful for human UGT or DDOST genes
Sequence 21 BP; 7 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                Determination of enzymes participating in glucuronic acid human being, comprises use of oligonucleotide probes.
                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-378271/41
                                                                                                                                                                                                                                                                                                                                                                                                                      (SAKA ) OTSUKA SEIYAKU
                                                                                                                                                                                                                                                                                                              being,
                                                                                                                                                                                                                                                                         8; Page 13; 13pp;
                                                                                                                                                                                                                                                                                                                                                                                                                          KOGYO
                                                                                                                                                                                                                                                                           Japanese.
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                                                                                                                                                                                                                                                                                                                                        conjugation
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Query Match
Best Local Similarity
Matches 17; Conserv
                     7250 TGGATGGGGAAATGTCTCTG 7269
20 TAGATGGGGGGATGTCTCTG
                                           Conservative
                                                       85.0%;
                                                                0.2%;
                                            0; Mismatches
                                                  Score 15.2; DB 1,
                                                                  DB 1;
                                                                 Length
                                             Indels
                                                                  21;
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07-OCT-2002
                                                                                                                                                     ABK88538 standard; DNA; 21 BP
                                                                                                      Homo sapiens
                                                                                                                   Panic disorder;
                                                                                                                            Human cholecystokinin associated PCR primer P2.
                                                                                                                                             ABK88538
                                                                                                                                     (first entry
                                                                                                                   polymorphism; human cholecystokinin; upper stream;
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JP2002171990-A.

18-JUN-2002.

08-DEC-2000; 2000JP-00375090

08-DEC-2000; 2000JP-00375090

(RIKA) RIKAGAKU KENKYUSHO.

WPI; 2002-569888/61.

Diagnosis and identification of panic disorder caused by polymorphism of upper stream region of human cholecystokinin gene.

Claim 6; Page 6; 13pp; Japanese.

The invention describes a method of diagnosing a panic disorder with polymorphism of the upper stream region of human cholecystokinin (CCK gene. This sequence represents a human cholecystokinin gene associate PCR primer associated

Sequence 21 BP; σ ₽ 5 C; 4 G; 6 T; 0 U; 0 Other,

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Query Match
Best Local Similarity
Matches 17; Conserv
Conservative
         0.2%;
Score 15.2; D
Pred. No. 1.8e
0; Mismatches
 0
         .8e+03;
                 DB 1;
                 Length
 Indels
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RESULT 2592
ABK16848/c
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                                          RESULT 2593
ABZ31311
                                                                                                                                                          Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                           The invention relates to a host cell, made to express an insoluble or aggregated protein having free cysteines residues. The cell is then lysed by chemical, enzymatic or physical agents and solubilised by exposing it to a denaturing agent, a reducing agent and a cysteine blocking agent, and is refolded into a biologically active form by reducing the concentrations of denaturing and reducing agents. The protein may belong to the growth hormone supergene family or may be an anti-angiogenesis factor. The method is useful for preparing a refolded, soluble form of an insoluble or aggregated protein. The proteins of the invention can act as delivery vehicles for enhancement of the circulatory half-life of the therapeutics that are attached or for directing delivery of a specific target within the body. Sequences ABK16774-ABK16852 represent PCR primers used in synthesis of the proteins
ABZ31311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Making and refolding insoluble or aggregated proteins having free cysteine by exposing host cell expressing protein to cysteine blo agent, and exposing to cysteine reactive group to increase their
                         ABZ31311 standard; DNA; 21 BP
                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 14; Page 52; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-089843/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein refolding; growth hormone supergene family; human; mouse; therapeutic half-life; PCR primer; anti-angiogenesis factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human protein refolding PCR primer #71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-MAY-2000; 2000US-0204617P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BOLD-) BOLDER BIOTECHNOLOGY INC.
                                                                                                                              6435
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                                                                                                   21
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                                                                                                                                                                         Similarity
                                                                                                                                                                                                                    21
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                                                                                                   ATTCATCTCAGCAGCAGTGT 2
                                                                                                                              ATTTAGCTAAGCAGCAGTGT 6454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NS,
                                                                                                                                                                                                                    BP; 6
                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001WO-US016088
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                                                                                                                                                                                                                    A; 4 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                        0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Doherty DH;
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                                                                                                                                                                                                                    G; 5 T; 0
                                                                                                                                                                        Score 15.2;
Pred. No. 1.
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                                                                                                                                                           Mismatches
                                                                                                                                                                                                                    U; 0 Other;
                                                                                                                                                                         1.8e+03
                                                                                                                                                                                       DB 1; Length 21;
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                                                                                                                                                          Gaps
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RESULT 2594 ABS98398/c ID ABS9839 XX

ABS98398 standard; DNA;

21 ВP 문 S

N

Best Loc Matches

Local

l Similarity

0,

Mismatches No.

Indels

0

Gaps

0

Pred.

..8e+03;

5695

CTGTTTTGCCTTCCTTTTCC 5714

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CC calls in which both alleles of a gene are modified, comprising modifying CC calls in which both alleles of a gene are modified, comprising modifying CC one allele by insertion or replacement by a cassette having an CC expressible selectable marker and modifying other allele by CC promoter, so that expression of the second allele is regulated by the CC promoter, which both alleles of a gene are modified. The diploid fungal CC cells in which both alleles modified are useful for identifying a gene that CC is essential to the survival or growth of a fungus, a gene that CC is essential to the virulence and/or pathogenicity of a fungus, a gene that contributes to the virulence of a diploid fungus to an antifungal agent that that inhibits the growth of a diploid fungus a gene condisease. (M1) is useful for identifying a compound which modulates the CC disease. (M1) is useful for identifying a compound which modulates the CC compound catabolism, biosynthetic, transporter, transcriptional, compound catabolism, biosynthetic, compound catabolism, biosynthetic, catabolism, bio
                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                         Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-DEC-2000; 2000US-0259128F
20-FEB-2001; 2001US-00792024
22-AUG-2001; 2001US-0314050F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Candida albicans GRACE strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roemer T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-DEC-2001; 2001WO-US049486
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002-566694/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 5530; 167pp + Sequence Listing; English.
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                                                                       B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                       A; 7 C; 4 G; 10 T; 0 U; 0 Other;
0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Boone
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                        Score 15.2; DB 1; Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bussey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ohlsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5530
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CC This invention relates to the sequence of an isolated nucleic acid CC molecule comprising at least one base variation from that of a known CC human cytochrome P450 A2 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), CC cytochrome P450 A2 (CYP4501A1), CC cytochrome CYTOX2, cytochrome P450 A2 (CYP4501A1), CC cytochrome CYTOX2, cytochrome CYTOX2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Whuman; ds; cytochrome P450 Al; CYP4501Al; UGT2B4; MDR1;

KW cytochrome P450 A2; CYP4501A2, cytochrome P450 02E; CYP45002E1; LTF;

KW cytochrome P450 A2; CYP4501A2, cytochrome P450 02E; CYP45002E1; LTF;

KW cytochrome P450 A2; CYP4501A2; aryl hydrocarbon; AHR; MRP3; NR112;

KW cytochrome P450 A2; CYP4501A2; aryl hydrocarbon; AHR; MRP3; NR112;

KW cytochrome P450 A2; CYP42; Aryl hydrocarbon; AHR; MRP3; NR112;

KW cytochrome P450 A2; CYP42; Aryl hydrocarbon; AHR; MRP3; NR12;

KW cytochrome P450 A2; CYP42; S-lipoxygenase activating protein; FLAP;

KW glutathione-S-transferase 12; GST12; histamine-N-methyl transferase;

KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;

KW NADPH quinone oxidoreductase 2; NQO2; sulfortansferase thermolabile; STM;

KW MADPH quinone oxidoreductase 2B4; UDP-glucuronosyl transferase 2B7;

KW WDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7;

KW WUP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;

KW multidrug resistance 1; lactotransferrin; orphan nuclear receptor;

KW multidrug resistance associated protein 3; cancer; prostate;

KW multidrug resistance associated protein 3; cancer; prostate;

KW multidrug resistance associated protein 3; cancer; prostate;

KW acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;

KW acetylcholine muscarinic receptor; prostate;

KW central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human multidrug resistance associated protein 3 polymorphic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABS98398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-698522/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Guida M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-NOV-2000; 2000US-00724389.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-NOV-2001; 2001WO-US044838.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 24; Page 152; 714pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder-related traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DNAS-) DNA SCI LAB INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens
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8888888888888888888
Matches
                 Query Match
Best Local
                                                                                                                                       metabolism. The polymorphic sequences contained in CYP4501A1, CYPP4501A2, AHR, MDR1 and/or MDR3 may also be used to screen individuals for susceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are used to screen for altered cardiovascular function, in COX2 for altered susceptibility to colorectal tunours, in DBI or CHMR1 for altered central nervous system function, in FLAP and HNMT for altered pulmonary, immunological or haematological function, in KLK2 for altered serine protease activity in the prostate, in LTF for altered fununological function, in CHMR3, CHMR4 or CHMR5 for altered central and peripheral nervous system function. The present sequence represents
                                                                               Sequence 21 BP; 3 A; 10 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                       polymorphic DNA sequence of the invention
17;
                        Similarity
Conservative
                 0.2%;
<u>.</u>
                                          Score 15.2;
                        Pred. No. 1.8e+03
  Mismatches
                                            DB 1; Length 21;
    ω
••
    Indels
  0,
    Сарв
      0
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RESULT 2595
ABK94358/c
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                                                                                            ABK94358 standard; DNA;
                                                                                     27-AUG-2002
                                                                                         ABK94358;
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                                                                                                      ATCCTTGGGGAAAGGGAGGA 1
                                                                                     (first entry)
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562 ATCCCTGGGGAAGGAAGGA 581

Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; 88. Synthetic Endothelin converting enzyme 1 (ECE-1) SNP detection primer #146

WO200224747-A2

28-MAR-2002

31-AUG-2001; 2001WO-EP010087

19-SEP-2000; 2000EP-00120123

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Brinkmann Ġ Hoffmeyer S

WPI; 2002-435060/46

Novel polynuclectide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the

Example 6; Page 67; 190pp; English

The invention describes a polynucleotide (I) of the endothelin (EDNN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. ((II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing a molecular variant gene comprising (I) is useful for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system

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RESULT 2596
ABK94357
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Best Local S
Matches 17
(EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) signaling system which is associated with a cardiovascular disease. (I), the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and polymucleotides encoding them are useful for preparation of a pharmaceutical composition for treating a cardiovascular disease such as coronary heart disease, hypertension, attherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polymucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polymucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding pathway cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
                                                                                                                                                                                           Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the
                                                                                                                                                                                                                                                                                                                                                                                                   31-AUG-2001; 2001WO-EP010087.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypertension; athérosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Endothelin
                                                                                                                                            Example 6; Page 67; 190pp; English
                                                                                                                                                                                                                                                                 WPI; 2002-435060/46.
                                                                                                                                                                                                                                                                                                                                                                   19-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200224747-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EDNR; signaling system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Endothelin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-AUG-2002
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17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCCCTCGATGTGGCCCAGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        converting
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                                                                                                                                                                                                                                                                                                                                                                   2000EP-00120123
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No. 1.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     enzyme; ECE; endothelin receptor; disease; coronary heart disease:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP detection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cc molecular variant gene comprising (I) is useful for identifying and cc obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system cor its gene product, or for identifying and obtaining an inhibitor of the CC Ticivity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and composition for treating a cardiovascular disease such as cc pharmaceutical composition for treating a cardiovascular disease such as cc pharmaceutical composition for treating a cardiovascular disease such as cc pharmaceutical angiogenesis or fatty acid metabolism e.g. diabetes and familial chypercholesterolaemia. The gene or a polynucleotide fragment of the CC pypercholesterolaemia. The gene or a polynuclocide markers, for cc creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or lost cells of the invention. This sequence represents a PCR primer used condentify single nucleotide polymorphisms in DNA encoding pathway cc cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP;
                                                                                                                               Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                    19-SEP-2000; 2000EP-00120123
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Endothelin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK94084;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EDN-1) SNP detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                   Hoffmeyer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphism;
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85.0%;
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Pred. No. 1.8e+03
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The invention desc (EDN)/endothelin a signaling system the gene encoding

tion describes a polynucleotide (I) of the othelin converting enzyme (ECE)/receptors system which is associated with a cardiov encoding EDN, ECE or EDNR (II) or a vector

cardiovascular disease.

expressing

of EDN endothelin

and

ECE

e. (I),

the

Example

9

Page 55; 190pp; English

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cc or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (CII), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing a cC molecular variant gene comprising (I) is useful for identifying and cC obtaining a pro-drug or drug capable of modulating the activity of a cell expressing a cC molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system cc or its gene product, or for identifying and obtaining an inhibitor of the cC activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and cC polymucleotides encoding them are useful for preparation of a polymuclar disease, hypertension, atherosclerosis, or related to coronary heart disease, hypertension, atherosclerosis, or related to hypercholesterolaemia. The gene or a polymucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for cc comprising polymucleotides, genes, vectors, polypeptides, antibodies or comprising polymucleotides, genes, vectors, polypeptides, antibodies or didntify single nucleotide polymorphisms in DNA encoding to cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway cc
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ABK94083/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK94083 standard; DNA; 21 BP
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                                                                                                     Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endothelin-1 (EDN-1) SNP detection PCR primer #27.
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                                                                                                                                                                                                                                                        WPI; 2002-435060/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
Example 6; Page 55; 190pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-SEP-2000; 2000EP-00120123
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                                                                                                                                                                                                                                                                                                                                                                                                   (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
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Pred. No. 1.8e+03;
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The invention describes a polynucleotide (I) of the endothelin

The present invention relates to a method for identifying compounds

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RESULT 2599
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Virucide; hepatotropic; antiinflammatory; antiviral; 2'-5'-oligoadenylate synthase; Flavivirus infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                        Claim 16;
                                                                                                     Identifying stimulators of oligoadenylate synas antiviral agents against Flavivirus, also
                                                                                                                                                                                                                Guenet J, l
Frenkiel M,
                                                                                                                                                                                                                                                                                                                                            04-APR-2001; 2001FR-00004598
                                                                                                                                                                                                                                                                                                                                                                                       04-APR-2002; 2002WO-FR001169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murine OAS gene isoform L1 PCR primer SEQ ID 13.
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                                                                                   for sensitivity to virus.
                                                                                                                                                                         WPI; 2003-058566/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3722 TCCTCATTCATTGAGCTTTTT 3742
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CNRS CENT NAT RECH SCI.
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                                      Page 80; 93pp; French.
                                                                                                                                                                                                                Mashimo T,
1, Despres
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                                                                                                                                                                                                                Simon-Chazottes D, Montagutelli X;
, Deubel V, Bonhomme F, Lucas M;
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lso mutated genes responsible
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PCR; primer;
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RESULT 2600
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Best Local S
Matches 17
The present invention describes the human brother of regulator of imprinted sites (BORIS) protein. Also described: (1) a vector comprising an isolated nucleic acid encoding BORIS; (2) a cell comprising the vector; (3) an isolated polypeptide molecule comprising a sequence encoding the human BORIS or its fragment comprising at least 307 or 21 contiguous amino acids, either one of which is optionally glycosylated, amidated, carboxylated, phosphorylated, seterified, N-acylated or converted into an acid addition salt and/or optionally dimerised or polymerised; (4) a cell line that produces a monoclonal antibody that is specific for a region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         that can stimulate a gene of the OAS (2'-5'-oligoadenylate synthase) family. The method comprises: (a) inducing expression of the OAS gene in a culture of cells from a non-human mammal (Flyr/Flyr or Flyr/Flyr; indicating resistance or sensitivity to Flavivirus infection); (b) treating cells with test compound; and (c) measuring activity of OAS gene relative to a control. (l) are potentially useful as antiviral agents for treating infections by Flaviviruses (e.g. hepatitis C; dengue; yellow fever and various forms of encephalitis). Genomic OAS DNA and derived cDNA, also the encoded proteins, are useful: (a) for treating Flavivirus infection; (b) in screening for anti-flavivirus agents, and (c) for evaluating sensitivity of subjects to Flavivirus infection and their likely response to interferon treatment, e.g. to identify patients at risk of developing severe forms of such infections. The present sequence is a PCR primer for murine OAS, which was used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-721780/68
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                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                          New nucleic acid molecule encoding a human or regulator of imprinted sites (BORIS), useful f
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse BORIS identification PCR
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ilarity 85.0%;
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Mismatches
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                                                                                                                                                                                                                                                                                                          composition
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The present invention relates to a method for the direct comparison numerical gene expression values between samples of genes. The methoc comprises amplifying cDNA in the presence of a competitive template inixture and primer pairs for several genes and then amplifying alique of the PCR products using a primer pair specific for each gene. The method of amplification is by multiplex standardised reverse transcriptase-polymerase chain reaction (StaRT-PCR). High density oligonucleotide or CDNA arrays are used to measure PCR products foll quantitative StaRT-PCR. The method is useful for the assessment of g

assessment of gene

following

Direct comparison of numerical gene expression values between samples of genes comprises using multiplex standardized reverse transcription-polymerase chain reaction.

Example

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NO 1; 59pp;

English

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the isolated polypeptide, where the region comprises any region that is recognisable by the monoclonal antibody other than one spanning a zinc finger region; (5) diagnosing cancer or predisposition to cancer in a male or female mammal; (6) predicting a predisposition to a cancer in an offspring of a male mammal; (7) prognosticating cancer in a mammal; (8) assessing the effectiveness of treatment of cancer in a mammal; (9) preventing or treating cancer that is due to the presence of BORIS nucleic acid or polypeptide; and (10) a composition comprising an inhibitor of BORIS and a carrier. BORIS has cytostatic activity, and can be used in gene therapy. A nucleic acid encoding BORIS can be used for preparing a composition for treating or preventing cancer. The present sequence represents a PCR primer corresponding to human BORIS and murine CTCF used to identify murine BORIS, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                          small
                                                                                                                                                                                                                                                (WILL/)
(CRAW/)
                                                                                                                                                                                                                                                                                                                                      02-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                        Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene expression;
StaRT-PCR; high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD56481 standard; DNA; 21
                                                                                                                                                                                                WPI; 2003-811730/76.
                                                                                                                                                                                                                       Willey JC,
                                                                                                                                                                                                                                                                                     28-MAR-2002; 2002US-00109349
                                                                                                                                                                                                                                                                                                             28-MAR-2002; 2002US-00109349
                                                                                                                                                                                                                                                                                                                                                               US2003186246-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7358
                                                                                                                                                                                                                                                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                               expression; multiplex standardised reverse transcriptase-PCR; -PCR; high density oligonucleotide array; cDNA array; biological sample; fine needle aspirate biopsy; captured microdissected material; human; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ш
                                                                                                                                                                                                                                               WILLEY J
CRAWFORD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTGTGAAATTATCCCAGCAG 7377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression
                                                                                                                                                                                                                        Crawford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 4 A; 4 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                ыO
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                                                                                                                                                                                                                        묩;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             analysis multiplex StaRT-PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 6 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15.2; DB 1;
No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              #1
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RESULT 2602
ADD90708
ID ADD9070
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                                                  Query Match
Best Local S
Matches 17
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression in small biological samples such as fine needle aspirate biopsites, and laser captured microdissected materials. The method allows for the standardised measurement of hundreds of genes from the same sample, which in prior art, could only be assessed for one gene. The present sequence represents a multiplex StaRT-PCR primer which can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                       Sequence
                                                                                                                              used for detecting the presence of allergies and for screening for treatments. The present sequence represents a PCR primer for mouse beta-
actin, which is used in an example from the present invention.
                                                                                                                                                                    detecting allergies; (2) method for screening for treatments for allergies; and (3) a kit for screening for treatments. The method can
                                                                                                                                                                                             The present invention describes a method for detecting an allergic disease using SOCS3 as a marker. The method comprises determining the level of expression of the marker gene in the sample and comparing it with that of a healthy individual. Also described: (1) a reagent for
                                                                                                                                                                                                                                                                                      Detecting allergic disease using SOCS3 as a marker
                                                                                                                                                                                                                                                                                                                                         Nagata N,
                                                                                                                                                                                                                                                                                                                                                                                                         27-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JAN-2003; 2003WO-JP000600.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse beta-actin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADD90708;
                                                                                                                                                                                                                                                               Example 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003072778-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   allergic disease; SOCS3; allergy; mouse; beta-actin; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD90708 standard;
                                                                                                                                                                                                                                                                                                                                                                    (NIGE-)
                                                                                                                                                                                                                                                                                                                                                                               (GENO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6802 CAGATTGGGAAGGAGGTATT 6821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
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                         527
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17; Conserv
 N
                                                   l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                    JAPAN
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                         CCATTGGCAACGAACGGGTC 546
                                                                                                       21 BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAGAATGGGGAGGAGATATT 1
 CTATTGGCAACGAGCGGTTC
                                                                                                                                                                                                                                                                                                                                         Oshida T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
ilarity 85.0%;
Conservative
                                                                                                                                                                                                                                                             SEQ ID NO 18; 80pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       method of the present invention.
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          2002JP-00052310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                    RES
GEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR
                                                                                                                                                                                                                                                                                                                                                                                 INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer
                                                                                                       5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                 85.0%;
                                                                                                                                                                                                                                                                                                                                         Sugita Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                                                   0
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Pred. No. 1.8e
0; Mismatches
                                                                 Score 15.2; DB 1
Pred. No. 1.8e+03
                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                           Kubo
                                                                                                                                                                                                                                                                                                                                           Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                           Saito
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                                                                           Length
                                                   Indels
                                                                                                                                                                                                                                                                                        for dermatitis
                                                                             21;
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RESULT 2604
AAQ78894
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Best Local
                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                               The sequences given in AAQ27805-7 are primers which are complementary to intronic sequences within the amyloid precusor protein gene (APP). They were used to amplify exon 17, APP encodes various isoforms which are precursors of beta-amyloid protein. The beta-amyloid protein is an approx. 4kD protein (39-42 amino acids) which is an internal cleavage product from APP. There are five distinct isoforms of APP containing 563, 595, 714, 751 and 770 amino acids. These are generated by alternative splicing of primary transcripts of APP which is located on human chromosome 21. The APP isoforms are glycosylated transmembrane proteins. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beta-amyloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
28-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ27806;
                                        Synthetic
                                                              25-MAR-2003
27-JUL-1995
                                                                                                                    AAQ78894 standard; DNA; 22
                                                                                                                                                                                                                                                             Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 31; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polynucleotide probe comprising nucleic acid encoding codon 717 of human APP770, useful for determining genetic pre-disposition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-284654/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hardy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UNLO ) IMPERIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-AUG-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9213069-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APP exon 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ27806 standard;
                    Hepatitis
                                                                                               AAQ78894;
                                                                                                                                                                                               5617 TTACCCAAGCTTCAAGGAAG 5636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JA,
                                                                                                                                                                          N
                                                                                                                                                                                                                               Similarity
                     W
                                        EcoRI-BglII fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chartier-Harlin MC,
                                                                                                                                                                                                                                                              BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                         disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer
                   virus preS2 gene; vaccine;
                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein;
                                                              (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91GB-00001307
91GB-00018445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92WO-GB000123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COLLEGE SCI TECHN MED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                             0.2%;
                                                                                                                                                                                                                                                              6 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amyloid precursor protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22
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                                                                                                                     ВP
                                                                                                                                                                                                                                                              e.
                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                             Score 15.2; DB 1
Pred. No. 1.9e+03
                                                                                                                                                                          21
                                                                                                                                                                                                                                                              3 T; 0 U;
                                          from plasmid p7,582-8.
                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Goate AM,
                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Owen MJ,
                                                                                                                                                                                                                                           1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         isoform;
                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mullan
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RESULT 2605
AAQ55161
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Best Local :
                                                                                                                                                                                                                                                                                                                                                               codons of preS2-S and a 4330 bp BandII-BandII fragment of vector pVAR15 comprising the vaccinia virus 7,5K promoter and thymidine kinase gene. The vector also carries the bla antibiotic resistance gene and is useful for production of a HBV vaccine. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                              virus, was constructed from a 1403 bp BamHI-BooRI fragment contg. the preS2-S gene (minus the first 3 N-terminal codons), an EcoRI-BglII synthetic linker fragment (AAQ708894) including the first 3 N-terminal codons of preS2-S and a 4330 bp BamHI-BamHI frament
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Recombinant plasmid p7,5S2-S, coding for preS2-S gene virus (HBV) - also methods for its construction and a virus expressing HBV preS2-S in mammalian cells.
                                WO9400569-A1
                                                     Synthetic.
                                                                                                             Sequence
                                                                                                                                   25-MAR-2003
22-JUL-1994
                                                                                                                                                                                           AAQ55161 standard;
                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Col 9; 5pp; Russian.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-364608/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Belyaev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SU1575576-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            misc_feature
                                                                                       Amyloid
                                                                                                                                                                    AAQ55161;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-JUN-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MOLE=) MOLECULAR BIOLOGY RES INST.
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                                                                                                                                                                                                                                                                                                           Similarity
                                                                                       precursor
                                                                                                             of
f
                                                                                                                                                                                                                                                                 TTGGATCTATCCTGATCTGG 4693
                                                                                                                                                                                                                                                                                                                                           22 BP; 8 A; 3 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                   TTAGATCTATCATGAACTGG
                                                                                                            PCR primer APP-17B for amyloid precursor protein
                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                 (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Putintseva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88SU-04445111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/label= EcoRI_sticky_end
/label= EcoRI_sticky_end
/note= "on the complementary strand, the 5'-end overhangs
the coding strand by the 4 base sequence 5'-AATT-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /codon_start= 14. .16
/note= "HBV preS2 translation initiation codon"
1. .4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                     protein; Huntington's chorea; Alzheimer's;
                                                                                                                                                                                           DNA;
                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                           22
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                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                        Score 15.2
Pred. No.
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                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                              for the preS2-S antigen of hepatitis bp BamHI-EcoRI fragment contg. the H
                                                                                                                                                                                                                                                                                                                      15.2;
                                                                                                                                                                                                                                                                                                          .9e+03;
                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of hepatitis B
strain of Vaccinia
                                                                                                                                                                                                                                                                                                                      22
                                                                                                                                                                                                                                                                                              0,
                                                                                   PCR primer;
                                                                                                            (APP).
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-JUN-1992;
07-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A yeast artificial chromosome (YAC) was prepd. that contd. the gene fo amyloid precursor protein (ABP). The APP YACK was introduced into embryonic stem (ES)cells (AB-1). For the identification of ES clones contg. APP sequences, individual DNA samples were pooled in rows and columns for PCR, and the pools analysed for APP sequences by PCR, usin primers APP-PA, APP-PB2, APP-17A and APP-17B, which are adapted from Pidani et al., Human Molecular Genetics, 1, 165-168, 1992. APP-PA/PB2 denote primers specific for the promoter region of the human APP gene, APP17A,17B are specific for exon 17. (Updated on 25-MAR-2003 to correct and application of the sequences)
                    WPI; 1995-275446/36.
                                           Pearse MJ,
                                                                                                    27-JAN-1994;
26-JAN-1995;
                                                                                                                                                                                                                                 Alpha-1,3-galactosyltransferase; alpha-1,3-GalT; transgenic hyperacute rejection; xerotransplantation; primer; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                     25-MAR-2003
10-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Transgenic non-human animals prodn., with via co-lipofection, provide useful models Huntington's chorea, Alzheimer's disease.
                                                                                                                                       27-JAN-1995;
                                                                                                                                                              03-AUG-1995.
                                                                                                                                                                                     WO9520661-A1
                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                               Alpha-1,3-GalT
                                                                                                                                                                                                                                                                                                                                        AAQ93086;
                                                                                                                                                                                                                                                                                                                                                              AAQ93086 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 8 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1994-026206/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Choi T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-JUN-1993;
                                                               (BRES-) BRESATEC LTD. (SVIN-) ST VINCENT'S HOSPITAL MELBOURNE LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENP-)
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 33; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Loring JF,
                                           Crawford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                              polyA primer
                                                                                                   94US-00188607
95US-00378617
                                                                                                                                       95WO-IB000088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-00900972.
93US-00001493.
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                                           Ŗ,
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                                           Robbins
                                                                                                                                                                                                                                                                               (reverse)
                                                                                                                                                                                                                                                                                                                                                               ВP
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                           Ŗ
                                           Rathjen
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                                           AJF;
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diseases e
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New alpha-1,3-galactosyltransferase and leukaemia inhibitor factor

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AAT78987/c
                                                                  Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8987/c
AAT78987
                                                                                                                                   This oligonucleotide comprises the 5' donor site of intron 2 of the mouse Huntington's disease (HD) gene (see also AAT78974). The splice site equences for the first 5 exons of the mouse and human HD gene were compared (see AAT78985-T79002). Targeted disruption of the murine HD gene, e.g. at exon 5, can be used to examine the function of the HD gene and its role in development. Transgenic mice can be used as models of HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The primers given in AAQ9308 and cattle alpha-GalT genes GalT cDNA (AAQ93077) by PCR correct'PR field.)
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gene; for
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Huntington's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse Huntington's disease gene intron 2 5' donor site.
                                                                                                                                                                                                                                                                                                                       (UYBR-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4459 IGGACITITITITITITIT 4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                          screening;
                                                                                                                                                                                                                                       Huntington's Disease gene - useful for model of Huntington's Disease.
                       22
                                                                   l Similarity
17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Page
                                       TTTTTTGTAAAAGATAATT 5493
                                                                                                                22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA and nucleic acid constructs for inactivating the transferase reliminating hyperacute region in human transplants.
                                                                                                                BP;
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease; animal model; transgenic animal; mouse; therapy;
ng; Hdh gene; ss.
                                                                                                                                                                                                                                                                                                                       BRITISH COLUMBIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ93081-86 are based on conserved regions T genes and were used to isolate porcine alpoy by PCR amplification. (Updated on 25-MAR-20
                                                                             0.2%;
                                                                                                                                                                                                                   69pp; English.
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                                                                             Score 15.2; DB 1;
Pred. No. 1.9e+03;
                                                                                                               G; 8 T; 0 U; 0 Other;
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                                                                   Mismatches
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                                                                                        DB 1;
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                                                                                        Length
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                                                                                                                                                                                                                                                     transgenic
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RESULT 2609
AAV99612
ID AAV9961
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RESULT 2608
RAX78249/c
ID AAX7824
XX AAX7824
XX AAX7824
XX Puman c
XX Puman c
XX Cyclin
KW Cyclin
KW Transec
XX Synthet
OS Homo se
XX I1-DEC
                                                                                                                                                                                                                                                                                                                               This invention describes a novel human transcriptional co-activator, CC designated cyclin T1 (also known as cyclin K), which interacts with the chuman immunodeficiency virus (HIV) Tat protein. Cyclin T1 is capable of participating as a constituent of the Tat-associated kinase ('TAK): Transcription elongation factor submit b (TEFP) complex. The polyapptide modulates Tat transactivation by enhancing the affinity of the Tat protein with the cis-acting transactivation response element ('TAR) RNA. Compounds identified by methods of the invention that disrupt the association of divalent cation metal(s) with cyclin T1 and/or Tat C protein are useful for the modulation of Tat transactivation in C biological systems. Cyclin T1, or antibodies spainst cyclin T1 can be used to treat a subject infected with HIV. Antibodies spainst cyclin T1 can be used as immunogens and also for studying tissue localization of protein C and complexes, for purification of inhibitors and are useful for c diagnostic applications. The probes and primers are useful for c identification and amplification of the cyclin T1 DNA. The transgenic calisate for tity infection
                                                                                                                     Best Local Similarity Matches 17; Conserv
                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cyclin T1; cyclin K; human; TAT protein; transcriptional coactivator; human immunodeficiency virus; HIV; Tat-associated kinase; TAK; TEFb; Transcription elongation factor subunit b; TAK/TEFb complex; cis-acting; transactivation response element; TAR; divalent cation metal; modulator; treatment; infection; immunogen; tissue localisation; diagnosis; transgenic animal; PCR primer; ss.
                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-1998;
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                                                                                                                                                                                                                                                                                                                  VIH
                                                       CTGTTTTGCCTTCCTTTTCC 5714
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              a transcriptional co-activator that interacts with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wei
                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                  infection
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98US-00126980.
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                                                                                                                                                                                                                                                 A; 4 C; 6
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                                                                                                                                                     0.2%;
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                                                                                                                        Score 15.2; D
Pred. No. 1.9e
0; Mismatches
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                                                                                                                                                                                                                                                 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                         .9e+03
                                                                                                                                                                                         DB 1;
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AAV99612 standard; DNA; 22

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AAX26493/c
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Best Local :
                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                            This is the nucleotide sequence of maize clpP gene primer clpP#3. The 5' nucleotide of the primer corresponds to nucleotide 70549 of the complementary strand of the maize plastid genome sequence. The primer was designed to add a XhoI restriction site downstream of an amplified clpP fragment following PCR amplification. The PCR product was cloned into vector pBSKS+ and used to generate protecting RNA for use in in vitro capping experiments. The invention provides isolated rpoB, atpB, clpP and 16S rDNA NEP and PEP promoter elements (see AAV99569-99) useful for producing exogenous proteins of interest in plant plastids
                                                                                                                                                                                                                                                                                                                                         Sequence
18-FEB-1999
                     WO9907890-A1
                                                                                                       cDNA transcript resulting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 17; 79pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated nuclear-encoded plastid RNA polymerase promoter sequences useful for expressing exogenous protein in plant plastids such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-JUN-1997;
12-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transgenic
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                                             Synthetic
                                                                    selection oligonucleotide;
                                                                                  Isolation;
                                                                                                                                 26-MAY-1999
                                                                                                                                                         AAX26493;
                                                                                                                                                                                AAX26493 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-070262/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maliga P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Promoter; nuclear encoded plastid RNA polymerase; NEP; clpP; chloroplast; transgenic plant; maize; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV99612;
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                                                                                                                                                                                                                                                                                                      Loca1
                                                                                                                                                                                                                                                             3342 GAATCCAGTTTGTAGGAAGA 3361
                                                                                                                                                                                                                                           w
                                                                                                                                                                                                                                                                                          l Similarity
17; Conserv
                                                                                                                                                                                                                                           GAATTCTGTTTGTAAGAAGA 22
                                                                                                                                                                                                                                                                                                                                          22 BP; 7
                                                                    nucleic acid; subtraction oligonucleotide oligonucleotide; structure probing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Silhavy D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene
                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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97US-0058670P.
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                                                                                                                                                                                                                                                                                                                                          A; 1 C; 7
                                                                                                                                entry
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                                                                                                       from
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                                                                                                                                                                                                                                                                                                                                          G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         Score 15.2; D
Pred. No. 1.9e
0; Mismatches
                                                                                                       reverse transcription of the test RNA
                                                                                                                                                                                                                                                                                                      .9e+03;
                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                Length 22;
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AAZ32687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fragment from a mixture of nucleic acid fragments. The method comprises removing a non-target fragment by hybridising it to an immobilised "subtraction" oligonucleotide that is complementary to a sequence present in the non-target fragment but absent in the target fragment, repeating this step until a known sequence in the target nucleic acid is unique and then selecting the target fragment by hybridising to a complementary immobilised "selection" oligonucleotide. The method can be used in structure probing experiments to determine whether a given nucleotide is modified by a modifying agent, or whether a compound alters reactivity of a nucleotide in a test nucleic acid toward a modifying agent. The present sequence represents a cDNA transcript resulting from reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The
                                                                                                                                                                    US5981175-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-AUG-1998;
                                                                          07-JAN-1993;
18-JUN-1993;
                                                                                                                  25-JAN-1994;
                                                                                                                                            09-NOV-1999.
                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                      primer; ss.
                                                                                                                                                                                                                                                 APP; amyloid precursor protein; Alzheimer's disease; transfection; transgenic animal; animal model; disease; transgene; co-lipofection; yeast artificial chromosome; YAC; lipid; cationic; selectable; exon;
                                                                                                                                                                                                                                                                                                     Human APP
                                                                                                                                                                                                                                                                                                                                21-JAN-2000
                                                                                                                                                                                                                                                                                                                                                          AAZ32687;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Fig 9B; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid isolation, quantification and structural probing - using new hybridisation-based and ligation-dependEnt methods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stern
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-AUG-1997;
                                                 (GENP-)
                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                    AAZ32687 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transcription of the test RNA (AAX26474)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-167451/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYMA-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specification describes a method for isolating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5702 GCCTTCCTTTTCCTCTCTC 5721
                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
17; Conserv
                                                 GENPHARM INT INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCCTTCCCCTTCCTGTTCTC 2
                                                                                                                                                                                                                                                                                                       exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MASSACHUSETTS.
                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                          93US-00001493
93US-00079444
                                                                                                                  94US-00187161.
                                                                                                                                                                                                                                                                                                     17-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98WO-US016211
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
                                                                                                                                                                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                    ₽₽
                                                                                                                                                                                                                                                                                                     PCR primer APP-17B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; DB 1;
Pred. No. 1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a target nucleic acid
The method comprises
                                                                                                                                                                                                                                                  selectable; exon;
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                                                                                                                                                                                                                                                    PCR;
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WPI; 1999-633306/54

Kay RM,

Choi T,

Loring

Production of transfected mammalian cells DNA species, useful for the production of

by co-lipofection with multiple transgenic animals for use as

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RESULT 2612
AAZ37259
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents human APP (amyloid precursor protein) exon 17-CC specific PCR primer APP-17B, used with PCR primer APP-17A (AAZ32686) to CC amplify exon 17 of the human APP gene in murine embryonic stem cells CC transfected via a novel method with a YAC (yeast artificial chromosome) CC containing the human APP gene. The novel method of transfection produces CC a selectable co-lipofected mammalian cell incorporating multiple CC teterologous DNA species. It comprises forming a co-lipofection complex CC comprising a cationic lipid, a first polynucleotide larger than 50 kb, CC and an unlinked second polynucleotide comprising a selectable marker gene CC expression cassette, and transfecting mammalian cells with it. Both CC expression cassette, and transfecting mammalian cells with it. Both CC electable co-lipofected mammalian cells which contain incorporated CC multiple heterologous DNA species. The method can be used for transferring large segments of DNA, such as large YAC clones, into CC mammalian cells such as embryonic stem cells. The methods can be used for transgenic animals as models for Alzheimer's disease. The methods can be used to produce transgenic animals as models for Alzheimer's disease. The methods can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                   Synthetic.
Gallus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA species, useful for disease models.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Col
                            New polypeptide, antibodies.
                                                                                                                                                                                                                                                                                                                                               AV37 antigen; monoclonal antibody; hybridoma AV37; vaccine; avian tumour; oncogenic avian virus; Marek's disease virus; avian leucosis virus; Rous-associated virus; reticuloendotheliosis virus; therapy; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                    Burgess SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ37259 standard; DNA;
                                                                        WPI; 2000-013437/01
                                                                                                                                                                 29-APR-1998;
                                                                                                                                                                                               22-APR-1999;
                                                                                                                                                                                                                            04-NOV-1999.
                                                                                                                                                                                                                                                         WO9955860-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer for AV37 antigen coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JAN-2000
                                                                                                                                   (ANIM-) INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5617 TTACCCAAGCTTCAAGGAAG 5636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TAACCCAAGCATCATGGAAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                   ANIMAL HEALTH LTD
                                                                                                       Davison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                 98GB-00009070.
                                                                                                                                                                                               99WO-GB001238
                                           useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                       TF,
                                             8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22 BP
                                                                                                        Ross LJN;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.2; DB 1;
Pred. No. 1.9e+03;
0; Mismatches 3
                                             vaccine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                           generate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 22;
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                                               monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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Claim 31; Page 39; 63pp; English

RESULT 2614 AAA39764 ID AAA3976

AAA39764 standard; DNA; 22

BP

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AGCCAGAGCAGCAGCAG 20

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RESULT 2613
AAC61082
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Best Local
                                          Query Match
Best Local (
                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence is a PCR primer for DNA encoding the AV37 antigen protein of the invention. The protein is recognised by a monoclonal antibody (MAb) secreted by the hybridoma AV37 deposited at the European Collection of Cell Cultures (ECACC) accession number 98030304. The polypeptide can be used to isolate a MAb, produce a hybridoma producing the MAb, and in a composition for use as a vaccine a hybridoma producing the MAb, and in a composition for use as a vaccine. The vaccine can be used against oncogenic avian viruses, including Marek's disease virus, avian leucosis virus. Rous-associated virus and reticuloendotheliosis virus. The vector
                                                                                                           includes DNA and protein sequences for the receptor (AAC61078 and AX885916). Also included in the invention is an antibody specific for the murine nurse cell receptor. DNA encoding the receptor can be used to detect Digeorge's syndrome. The present sequence represents a primer used in the isolation and characterisation of the receptor of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   can be used to treat avian tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC61082;
                                                                                   Sequence 22 BP; 7 A; 7 C; 7 G; 1 T; 0 U;
                                                                                                                                                                                 This invention relates to a mouse nurse cell receptor.
                                                                                                                                                                                                                Claim 9; Page 7; 27pp; Japanese.
                                                                                                                                                                                                                                          A mouse nurse cell receptor gene
                                                                                                                                                                                                                                                                                                                              24-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                        24-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                JP2000236882-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse; nurse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer B6K311 used in isolation of cDNA encoding nurse cell receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC61082 standard; DNA;
                                                                                                                                                                                                                                                                                                  (SHIO ) SHIONOGI & CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5734 TTCCTTTCCCTTTTCTA 5753
  7405 AGCANCATCAGCAGCAGCAG 7424
                             cal Similarity
17; Conserv
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17; Conserv
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                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                               99JP-00046603
                                                                                                                                                                                                                                                                                                                                                          99JP-00046603.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 11 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
                                           0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detection; DiGeorge's
                             0;
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Pred.
                                             Score 15.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15.2;
                                           .9e+03
                                                                                      0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        syndrome; primer; ss
                                Indels
                                                                                                                                                                                       The invention
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                                0
                                Gaps
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RESULT 2615
AAZ94181/c
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  2222222222223B
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Best Local
                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel heat-inducible promoter (1). (1) is used to express foreign genes in fungi, particularly yeast and specifically Hansenula polymorpha, especially proteins that require glycosylation and can not be expressed in bacteria. (1) is active even at high temperature, allowing yeast to be cultured at higher than normal temperatures with reduced contamination by other microorganisms and reduced need for expensive cooling. (1) is selectively activated by heat (it contains no general stress-responsive elements), so permits expression of foreign genes to be suppressed during selected stages of growth, resulting in less injury to cells and biomass. This sequence represents a primer used in the isolation of the trehalose-6-phosphate synthase TPS1 gene from Hansenula polymorpha. (Updated on 15-SEP-2003 to standardise OS field)
           GABAB receptor 2; GABAB-R2; human; bladder disorder; gastrointestinal disorder; central nervous system disorder; lung disorder; spasticity; epilepsy; Alzheimer's disease; pain affective disorder; feeding disorder; diagnosis; gene therapy; G-protein coupled receptor; GABA; gamma-aminobutyric acid;
                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression of foreign genes in yeast, contains no general
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pichia angusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Trehalose-6-phoshate synthase; TPS1; heat shock inducible;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-SEP-2003
18-SEP-2000
                                                                                               Human
                                                                                                                                                                              AAZ94181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RHEI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymorpha
                                                                                                                                                                                                                                                                          7190
                                                                                            GABAB-Rla PCR primer
                                                                                                                                                                                                                                                                                                    17; Conserv
                                                                                                                                                                                                                                                 w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RHEINBIOTECH GES NEUE BIOTECHNOLOGISCHE.
                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                               GTGTGACTACTCTGGTTTTC 7209
                                                                                                                                                                                                                                                                                                                                                        22 BP; 2 A; 3 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                               GTGTGATTACTGTGGTTTGC 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           De
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 12; 19pp;
                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TPS1 DNA primer F9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99CH-00000279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99CH-00000279
                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                              0.2%;
85.0%;
PCR primer; GABAB-Rla;
                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 German.
                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                 Score 15.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                 .9e+03;
                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for controlling stress response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                promoter;
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is that of a primer used in a PCR-RACE in order t generate a 800 bp fragment of GABAB receptor GABAB-R1a CDNA. The invention relates to novel GABAB subtypes GABAB-R1c and GABAB-R2 (see AAY79202). It also relates to variants of the receptors, nucleotide sequences (see AAZ94168) encoding the receptors, vectors, stable cell lines, antibodies, screening methods of receptor production, methods of treatment or prophylaxis of a disorder that is responsive t modulation of GABAB receptor activity. The disorder is especially a central nervous system disorder, a gastrointestinal disorder, a lung disorder or a bladder disorder, especially spasticity, epilepsy, alzheimer's disease, pain or an affective or feeding disorder (claimed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GABA-B receptor subtypes useful for identifying modulators of GABA-B receptor activity that may be used for preventing and treating diseases including Alzheimer's disease, epilepsy and spasticity.
(KYOW ) KYOWA HAKKO KOGYO KK
                                                                            26-OCT-2000.
                                                                                                                                                                 Human; histidine domain-protein tyrosine phosphatase; HD-PTP; chromosome 3p21.3; gene deletion; tumour suppressor; cytostat lung cancer; tumour; gene therapy; diagnosis; recombinant pro
                                                                                                                                                                                                                                                                                              AAC81233 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-SEP-1998;
09-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                  14-APR-2000;
                                                                                                   WO200063392-A1
                                                                                                                                                                                                                    Human tyrosine
                                                                                                                                                                                                                                              23-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 29; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barnes AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 3 A; 5 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                  2000WO-JP002455
                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                   phosphatase HD-PTP exon
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98US-0103670P
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                                                                                                                                                       primer; ss
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                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.2; DB 1
Pred. No. 1.9e+03
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                                                                                                                                                                                                                    2 PCR
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                                                                                                                                                                                                                    primer,
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RESULT 2617
AAC80270/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          domain-protein tyrosine phosphatase (HD-PTP; AAB29661) and to human HD-PTP protein tyrosine phosphatase (HD-PTP; AAB29661) and to human HD-PTP nucleic acids (AAC81224, AAC81225, AAC81262). The HD-PTP CG gene is located on chromosome 3p21.3. This region is frequently found to be deleted in lung cancers, and is therefore thought to contain a tumour suppressor gene. The invention also relates to expression vectors and host cells containing human HD-PTP nucleic acids; the recombinant production of HD-PTP; anticancer drugs containing HD-PTP; gene therapy compositions containing DNA encoding HD-PTP; diagnostic reagents containing HD-PTP pligonuclectides; antibodies specific for use in cancer diagnosis and investigation. HD-PTP specific antibodies for use in cancer diagnosis and investigation. HD-PTP proteins, nucleic acids and cancers, particularly those of the lung. The present sequence represents a human HD-PTP PCR primer used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 17; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLA-A; HLA-B; HLA-C; typing; primer; human; ss.
                                                                                                                                                                                                                                                                                                                                                         09-APR-1999;
11-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reverse primer #98 used for amplification of HLA-A exon
                                                                                                                                                                              WPI; 2000-647426/62.
                                                                                                                                                                                                                                                                                               (INNO-) INNOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                           05-APR-2000;
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                                                                                                             Locus-specific,
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                                                                                                                                                                                                                                  I, Rombout
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 7 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
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99US-0138614P.
                                                                                     separate amplification of exon 2, exon 3, ar
yte antigen (HLA)-A, HLA-B, or HLA-C alleles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
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to human HD-
. The HD-PTP
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Claim

4; Page 40; 128pp; English

of human leukocyte antigen primer sets, useful for sub

subtyping

or typing of HLA Class I alleles.

defined

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CC with primer sequence is that of primer sequence, which was used CC with primer sequence is that of primer sequence with primer sequence is that of primer sequence which was used CC with primer sequenced for the PCR amplification of CC expression patterns of sequenced to determine relative CC expression in healthy lung, with very little expression in healthy CC prostate, high levels of expression in liver and pancreatic carcinomas and moderate levels in 2 of 6 prostate, kidney and ovarian carcinomas, and CC at moderate levels in 2 of 6 prostate cancer samples. The 25 psg CC polynucleotides of the invention (see AAH26845-69) are diagnostic markers CC host cells, PSG polypeptides, and antibodies immunospecific for the CC polypeptides are claimed. Also claimed are methods and tools for using CC the markers for diagnosing prostate cancer, diagnosing metastasis of CC prostate cancer, staging prostate cancer, diagnosing prostate cancer, continuing prostate cancer, the context of the context of the context of the context of the cancer of the context of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel prostate cancer specific genes and polypeptides encoded by the genes, useful for detecting, diagnosing, monitoring, staging, prognosticating, imaging and treating prostate cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 56; 83pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; prostate cancer; diagnosis; therapy; imaging; vaccine; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human prostate specific gene clone sqpro030 reverse PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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Pred. No. 1.9e+03
0; Mismatches
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                                                                                                                       CC designated BGTNC#10 (AAB60393) and to the CDNA clone encoding it

CC (AAF27486). The human BGTNC#10 gene is located on chromosome 22q11, in a

CC region associated with DiGeorge's syndrome (a congenital disorder which

CC is often associated with heart defects, anomalies of the major vessels,

CC oesophageal atresia and abnormalities of facial structures). The human

CC BGTNC#10 gene was identified in a human genomic sequence (GenBank

CC (AC005500) via homology with a murine BGTNC. Appropriate PCR primers

CC (AAF27479-AAF27485) were designed to isolate the human BGTNC#10 CDNA from

CC (AAF27479-AAF27485) were designed to isolate the human BGTNC#10 CDNA from

CC (AAF27479-AAF27485) were designed to isolate to expression vectors

CC (AAF27479-AAF27485) were designed to isolate the human BGTNC#10 CDNA; the recombinant

CC and transformants comprising the human BGTNC#10; a monoclonal antibody

CC recognising BGTNC#10; a hybridoma producing such an antibody; a method of

CC compounds thus identified. The human BGTNC#10; a monoclonal antibody

CC screening for a ligand which binds to a human nurse cell receptor; and

CC compounds thus identified. The human BGTNC#10 CDNA can be used in the

CC detection and diagnosis of DiGeorge's syndrome. The present sequence

CC represents a mouse nurse cell receptor BGTNC#10 RACE (rapid amplification

CC of CDNA ends) PCR primer used in an exemplification of the invention
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Best Local Similarity
Matches 17; Conserv
                                                                 Query Match
Best Local
                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nurse cell receptor; mouse; murine; B6TNC#10; DiGeorge's syndrome; drug screening; detection; diagnosis; rapid amplification of cDNA ends;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 8; 48pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide which is a human nurse cell receptor is useful for screening assays, especially for DiGeorge's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse nurse cell receptor B6TNC#10 RACE PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF27472 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a human nurse cell receptor protein sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-161956/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-FEB-2000; 2000JP-00045321
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4688
7405 AGCAACATCAGCAGCAGCAG 7424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                   1 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22
                                                                                                                     22 BP;
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nilarity 85.0%;
Conservative
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llarity 85.0%;
Conservative
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                                                                                                                       7
                                                                                                                       Ą
                                                                                                                       7 C; 7 G; 1 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 C; 3 G; 7 T; 0 U; 0 Other;
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                                                                   Score 15.2;
Pred. No. 1.
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Pred. No. 1
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                                                                                                                       0
U;
                                                                     1.9e+03;
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                                                                                                                       0 Other;
                                                                                     DB 1;
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                                                                                 Length 22;
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                                                   Indels
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                                                   Gaps
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                                                                Matches
                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                           mRNA-protein (mRNP) complexes. The method comprises contacting a biological sample comprising mRNP complex with ligand that specifically binds a component of mRNP complex, separating mRNP complex by binding the ligand with a molecule specific for ligand, which is attached to the solid support and then collecting the mRNP complex by removing the complex from the support. The method is useful for in vivo partitioning of cellular mRNA protein complexes in a biological sample. The method is useful for determining the ribonomic profile of a cell which has numerous uses including monitoring of tumour development, state of growth or state of development, perturbations of a biological system such as disease, drug or toxin treatment and the state of cell aging or death, attributed the state of cell aging or death.
                                                                                                                                                                           distinguishing ribonomic profiles among organisms, to discriminate between transcriptional and post-transcriptional contributions to gene expression and to track the movement of RNAs through RNP complexes, including the interactions of combinations of proteins with RNAs in RNP complexes. AAH28281-AAH28316 represent sequences derived from the 3' untranslated region (UTR) of mRNA of various genes. The sequences contains
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 6; Page 31; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Partitioning endogenous mRNA-protein complexes in vivo, by contacting sample comprising the complex with ligand that binds to a component of the complex and separating complex by binding ligand with a binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA protein complex; tumour development; cell ribonomic profile; RNA-binding protein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3' untranslated
                                                                                                                                                             target sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The specification describes a method for partitioning endogenous cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Keene JD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH28299 standard; RNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (KEEN/) KEENE J D.
                             4020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-425706/45
22
                            AAAAAAGAGAGAAAACAAAA 4039
                                                                                                                               22 BP; 2
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                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    region sequence from neuronal cadherin
                                                                                                                                                             for RNA-binding proteins
                                                                                                                              A; 1 C; 1 G; 0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SA,
                                                                              0.2%;
                                                               0;
                                                                             Score 15.2; DB 1;
Pred. No. 1.9e+03;
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                                                               Mismatches
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                                                                                                                               18 U;
                                                                                              DB 1;
                                                                                                                               0 Other
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                                                                                            Length 22;
                                                               Indels
                                                               0;
                                                               Gaps
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RESULT 2621 AAH28297/c

ID _ AAH28297

standard; RNA; 22

BP.

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BXGXAXB
                                                                                                         RESULT 2622
ABL35693
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                                                                                                                                                                                                                                                                                                                      CC mRNA protein (mRNP) complexes. The method comprises contacting a component of mRNP complex with ligand that specifically binds a component of mRNP complex with ligand that specifically complex by binding the condition of mRNP complex with ligand that specifically conditions a component of mRNP complex with ligand that specifically conditions are complexed to the condition of the support and then collecting the mRNP complex by removing the conditions of callular mRNA protein complexes in a biological sample. The method is useful for determining the ribonomic profile of a cell which has numerous conditions of a biological sample. The method is useful for determining the ribonomic profile of a cell which has numerous conditions of a biological system such as disease, and the conditions of a biological system such as disease, conditions of conditional conditi
                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mRNA protein complex; tumour development; cell aging; death; ribonomic profile; RNA-binding protein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Partitioning endogenous mRNA-protein complexes in vivo, by contacting sample comprising the complex with ligand that binds to a component of the complex and separating complex by binding ligand with a binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3' untranslated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-SEP-2001
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                                04-APR-2002
                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 31; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Keene JD, Tenenbaum
Immunostimulatory oligonucleotide SEQ ID NO: 619.
                                                                                            ABL35693
                                                                                                                                                                                                    4020 AAAAAAGAGAGAAAACAAAA 4039
                                                                                                                                                                       22
                                                                                                                                                                                                                                  l Similarity
17; Conserv
                                                                                            standard;
                                                                                                                                                                                                                                                                                               22 BP; 2 A; 1 C; 1 G; 0 T; 18 U; 0 Other;
                                                                                                                                                                         AAAAAATACAGAAAATAAAA 3
                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     region sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                describes a method for partitioning endogenous cellular
                                                                                                                                                                                                                                                  0.2%;
                                                                                              DNA;
                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carson
                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                  Score 15.2;
Pred. No. 1.
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                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                    .9e+03;
                                                                                                                                                                                                                                                                    DB 1;
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                                                                                                                                                                                                                                                                 Length 22;
                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                     0
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RESULT 2623
ABK97546
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Best Local
                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to an immunostimulatory composition, which comprises at least one oligonucleotide comprising both an RNA region and a DNA region. The composition is useful for enhancing an immune response or inducing cytokines. It can be used as a vaccine adjuvant and in treating diseases, including pathogenic infection, (non-)malignant cumours (e.g. cancers of the brain, lung, overy, breast, prostate or colon, or carcinomas and sarcomas), autoimmune diseases or allergies (e.g. allergic rhinitis, hay fever or food allergies). Lyme disease, preventing or ameliorating the symptoms resulting from exposure to a biovariare agent, e.g. Ebola, Anthrax or Listeria. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pna/rna hybrid; phosphorothioate backbone; immunostimulatory; vaccine; infection; allergy; cancer; hypersensitivity; bio-warfare; immunostimulant; antiallergic; cytostatic; antimorobial; anti-HIV; immunosuppressive; protozoacide; virucide; hepatotropic; gene therapy; antiinflammatory; antibacterial; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New immunostimulatory compositions comprising RNA/DNA hybrid oligonucleotides, useful for enhancing an immune response or cytokines, particularly for treating diseases, e.g. cancer, cytokines, particularly for treating diseases, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                         Sequence 22 BP; 1 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-2000; 2000US-0209797P
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ABK97546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BIOS-)
                                                                                                                                                                  5460 GTTCTTACTCTGATTTTTTT 5479
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                                                                                                                                                                                                                                                            Similarity
standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Flora M, Klinman DM;
                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "optionally thymidine is replaced by uracil form RNA or DNA/RNA hybrids. Thymidine is linked to least one other base through a ribose sugar"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
DNA;
                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                                                                                                                                            3 C; 2
  22
ВP
                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                            G; 16 T; 0
                                                                                                                                                                                                                                                                  Score 15.2; DB 1;
Pred. No. 1.9e+03;
                                                                                                                                 22
                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                               U; 0 Other;
                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                             22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or inducing
                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                         Gaps
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Lecithin-cholesterol acyltransferase; LCAT; Norum disease; gene therapy;

Human LCAT gene forward PCR primer #7

07-OCT-2002

(first entry)

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RESULT 2624
ABK40406
ID ABK4040
XX
AC ABK4040
XX
DT 15-JUL-
XX
DE Probe f
XX
KW Human;
KW leukaem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a new polynucleotide comprising a CC nucleotide sequence which is a polymorphic variant of a reference CC sequence for lecithin-cholesterol acyltransferase (LCAT). The invention CC is useful for identifying an association between a trait (preferably a CC clinical response to drug targeting LCAT) and at least one genotype or CC haplotype of LCAT gene. The method of the invention has applicability in Cdeveloping diagnostic tests and therapeutic treatments for Norum disease, CC fish-eye disease and atherosclerotic cardiovascular disease. The CC fish-eye disease and atherosclerotic cardiovascular disease. The CC diversity, anthropological lineage, the significance of diversity and CC lineage at the phenotypic level, paternity testing, forensic applications CC and for identifying association between the LCAT genetic variation and a CC trait such as level of drug response or susceptibility to disease. In CC didition, the methods for identifying the LCAT haplotypes present in CC individuals are useful in the development of drugs targeting LCAT. For CC example, determining the frequency of individual LCAT haplotypes in a CC trait of the development of drugs targeting LCAT and CC example, determining the frequency of individual LCAT haplotypes in a CC trait development of drugs targeting that are most the development of drugs targeting that are most constructed one of a collection (ABK97573) of PCR primers that CC were used in the methods of the invention to detect notwoenthisms in the construction of the produce of the development of the invention to detect notwoenthisms in the construction of the produce of the development of the invention to detect notwoenthisms in the construction to the construction to detect notwoenthisms in the construction to the construction to the construction to detect notwoenthisms in t
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     represents one of a collection (ABK97534-ABK97573) of PCR primers the were used in the methods of the invention to detect polymorphisms in human LCAT gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated polymorphic variant polynucleotide of lecithin-cholesterol acyltransferase gene, useful for studying expression and biological function of the gene, and for therapeutic, diagnostic or forensic
  Human; PRO; benign tumour; malignant tumour; lymphoid malignancy;
leukaemia; neuronal disorder; stromal disorder; blastocoelic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chew A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fish-eye disease; atherosclerotic cardiovascular disease; forensic; population diversity; anthropological lineage; paternity testing; human; polymorphism; PCR; primer; ss.
                                                                                                             15-JUL-2002
                                                                                                                                                                                                  ABK40406 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                          ABK40406;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-JAN-2001; 2001WO-US000092
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                                                                                                                                                                                                                                                                                                                                                    1849
                                                                   for
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                                                                                                                                                                                                                                                                                                                                                 CAGGTGAAGAACGTGGTCAA 1868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 BP; 6
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                                                                                                                                                                                                                                                                                                           CTGGTGCAGAACCTGGTCAA 21
                                                                     gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                             (first entry)
                                                                     amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;
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                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                     analysis
                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.2;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                     of human PRO7133.
                                                                                                                                                                                                                                                                                                                                                                                                                  1.9e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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RESULT 2625
AAS16493/c
ID AAS1649
XX
AC AAS1649
XX
DT 14-FEB-
XX
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밁 S

AAS16493;

(first entry)

AAS16493 standard; DNA; 22

ВP

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Matches
                                                                                                                   Query Match
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17-AUG-1999;
31-AUG-1999;
                                                                                                                                                                                          The present invention relates to the isolation of novel human PRO polypeptides (AAU86128-AAU86162) and the polypucleotide sequences encoding them. The PRO polypeptides, agonists, antagonists or anti-PRO antibodies are useful for treating benign or malignant tumours (e.g. renal, kidney, bladder, breast, etc), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic, glandular, macrophagal, stromal and blastocoelic disorder inflammatory, immune and angiogenic disorders. The polynucleotide sequences are also useful in gene therapy. The present sequence represents a probe used in the methods of the present invention
                                                                                                                                                           Sequence 22 BP; 0 A; 11 C; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thirty five nucleic acids encoding PRO polypeptides, useful for treating benign or malignant tumors, leukemias and lymphoid malignancies, inflammatory, angiogenic and immunologic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Marsters SA,
Watanabe CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-205567/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-NOV-1999;
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26-JUL-1999;
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11-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammatory disorder; immune disorder; angiogenic disorder; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               L5-SEP-1999;
                                                                                                Local
                                       6893 TGCTCTCCCTTACTCTACTC 6912
                                                                             l Similarity
17; Conserv
  N
TGCTCTCCCTTGCTCTTCCC 21
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                         Page 143; 302pp; English.
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Wood WI;
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99US-0149395P.
99US-0151689P.
99WO-US020111.
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Pan J, Pitti RM, Roy MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US028301
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                                                                                              0.2%;
                                                                             0; Mismatches
                                                                                        Score 15.2; DB 1;
                                                                                                                                                           G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gurney
Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hillan KJ;
Stone DM;
                                                                                                                     22;
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                                                                                                                                                                                                                                                               disorders
                                                                             Gaps
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RESULT 2626
ABL55000/c
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                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated functional Type II gonadotropin-releasing hormone receptor (Type II GnRH-R), or a peptide comprising at least a portion of exon I of Type II GnRH-R, nucleic acids encoding the receptor, an expression vector comprising the nucleic acid, a host cell transformed with the vector, a transgenic animal having the construct stably integrated into its genome, an antibody able to bind specifically to Type II GnRH-R. The Type II GnRH-R is useful in gene therapy. The Type II GnRH-R is particularly useful for inhibiting endogenous Type II GnRH binding to its native receptor in vivo or as a contraceptive. The receptor may also have roles in neural development and sexual arousal. The present sequence is a PCR primer used to isolate a nucleic acid encoding marmoset type II GnRH-R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-APR-2000;
17-JUN-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polypeptide, useful in gene therapy, as contraceptive or for inhibiting endogenous Type II GnRH binding to its native receptor in vivo, comprises Type II gonadotropin-releasing hormone receptor and polynucleotides encoding receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Callithrix jacchus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Marmoset; ss; type II gonadotrophin-releasing hormone receptor; GnRH-R; contraceptive; neural development; sexual arousal; gene therapy;
WO200213862-A2
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 24; 92pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transgenic animal; PCR primer; S1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Marmoset Type II GnRH-R PCR primer S1.
                                                                    immunosuppressive; dermatological; ant antidiabetic; antithyroid; autoimmune
                                                                                                                            Human lymphoma-specific immunoglobulin PCR primer VH1L.
                                                                                                                                                          08-OCT-2002
                          Homo sapiens.
                                                                                                  Human; lymphoma; immunoglobulin; B-cell mediated pathology; cytostatic;
                                                                                                                                                                                                                  ABL55000
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                                                                                                                                                                                                                                                                                                                   1976
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                                                                                                                                                                                                                                                                                                                                               17;
                                                        88.
                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      22
                                                                                                                                                                                                                                                                                                          CAGTGATATTCCTGGGAGCA 1995
                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                        CAGTGATATTCCAGGTGGCA
                                                                                                                                                                                                                                                                                                                                                                                                     B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lowe
                                                                                                                                                                                                                                                                                                                                              0.2%;
llarity 85.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000GB-00014761.
2000US-0215232P.
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                                                                                 dermatological; antiinflammatory; neuroprotective;
                                                                                                                                                                                                                  DNA;
                                                                                                                                                          entry)
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                                                                                                                                                                                                                                                                                                                                                                                                     6 C; 5
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                                                                                                                                                                                                                                                                                                                                                                                                     G; 5
                                                                                                                                                                                                                                                                                                                                              0,
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                                                                                                                                                                                                                                                                                                                                                           Score 15.2;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                     T; 0 U; 0 Other;
                                                                     disease; B-cell lymphoma;
                                                                                                                                                                                                                                                                                                                                                           .9e+03;
                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                       Length 22;
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RESULT 2627 ABS67627

ABS67627 standard; DNA; 22 BP

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20

CCAGGTCCAGTCCATGGTGA

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Matches

Conservative

0

Pred. No. 1.9 0; Mismatches

Indels

0

Gaps

0

7344 CCTTGTCCAGTCCATTGTGA 7363

Mouse casein 29-NOV-2002

kinase-2 RT-PCR primer

(first entry)

15-AUG-2002

Mus musculus tumour;

RT-PCR;

ss; antisense therapy; casein kinase-2 alpha; cytostatic; antidiabetic; antiinflammatory; diabetes; hyperproliferative disorder; cancer; PCR; breast cancer; prostate cancer; liver cancer; infection; inflammation;

reverse trancriptase PCR; primer;

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Colymphoma-specific immunoglobulin heavy and light chains. The invention of relates to a novel composition for altering a B-cell mediated pathology in a patient. The composition contains a chimeric protein comprising at Cleast a portion of a variable region of heavy chain or light chain (WH or CV L) linked to at least a portion of an immunoglobulin constant region, converted with a B cell clone from the patient chaving the B cell mediated pathology. The composition of the invention constant region, convergence of the pathology. The composition of the invention constant region, convergence of the pathology. The composition of the invention constant region, convergence of the pathology. The composition of the invention convergence of the pathology. The composition of the invention convergence of the pathology. The composition convergence of the pathology. This convergence is a vaccine useful for altering a B cell mediated pathology. This convergence is a vaccine useful for altering a B cell mediated pathology. This convergence is a vaccine useful for altering a B cell mediated pathology. This convergence is convergence of the pathology of the pathology is subscience, autoimmune thyroiditis and scleroderma. The pathology is convergence of the pathology is a cytokine e.g. granulocyte-macrophage-colony stimulating factor (GM-CSF) or chemokine e.g. monocyte chemotactic protein 3 (MCR 3)
  Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Composition for altering B-cell mediated pathology, has a chimeric protein having portion of variable region of heavy chain or light chain linked to portion constant region associated with patient B cell clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-AUG-2000; 2000US-0224722P
11-AUG-2000; 2000US-0224723P
23-MAR-2001; 2001US-0279079P
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                                                                     Sequence 22 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FAVRILLE INC
      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shopes RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 43; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 represents a PCR primer used in the invention to amplify cific immunoglobulin heavy and light chains. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001WO-US025204
                                                                     5 A; 6 C; 7
      0.2%;
                                                                       G; 4 T; 0 U; 0 Other;
Score 15.2; ub .
                              DB 1;
                         Length 22;
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RESULT 2628
ABL45313
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Best Local
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10-MAR-2000; 2000JP-00066716
                                 12-MAR-2001; 2001JP-00068285
                                                                                                                                                                                                                               Human chromosome 1p36-35 PCR primer SEQ ID NO:2357.
                                                                                                                                                                                                                                                                                                                                            ABL45313 standard; DNA; 22 BP
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                                                                                                       JP2001321190-A
                                                                                                                                         Homo sapiens.
                                                                                                                                                                                             Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                   11-APR-2002 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-JAN-2002; 2002WO-US002942.
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                                                                                                                                                                            primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2408 CCACAGTGGACACCAACATC 2427
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                                                                                                                                                                                                                                                                                                                                                                                                                                   CCACAGTGAAAACCAGCATC 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 2629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell and the cronssituted as the positions on the chromosome and arrayed. The order of the corresponding to the chromosome and arrayed. The antargate of the propersion of the chromosome and arrayed.
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(GENO-)
                                                                                                                       26-DEC-2000; 2000JP-00396167.
                                                                                                                                                                                      09-JUL-2002
                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                 PCR; primer; allergic disease; allergy; NB-1; E48 antigen; c-fos; involucrin; BENE; HBp17; fibronectin; Idl; ss.
                                                                                                                                                                                                                                                                                                                                 HBp17 reverse PCR primer
                                                                                                                                                                                                                                                                                                                                                                  31-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primers for human chromosome 1p36-35 DNA, and ABL45323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
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                                                                                                                                                      26-DEC-2000; 2000JP-00396167
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                                                                                                                                                                                                                                                                                                                                                                                                ABQ80518;
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                                                                                         (GENO-) GENOX SOYAKU KENKYUSHO
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GENOTEX YG.
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Inspection of allergic diseases, allergic diseases.

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The invention relates to a new fusion polypeptide or its pharmaceutically acceptable salt comprises at least one IgE-binding domain fused to at least one human serum albumin (HSA) component, where the IgE-binding domain is the sequence (a) defined residues Val26-Leu204 of the protein sequence appearing as ABG32801, or a truncation at the carboxy terminus by 1-12 amino acids. Also included are: (1) a fusion polypeptide defined by residues Val26-Leu278 of the protein appearing as ABG32803; (2) a polynucleotide sequence encoding the fusion protein; (3) a host cell transformed with the polynucleotide; (4) a method of preparing the fusion protein comprising transforming a host cell with a vector comprising transforming a host cell with a vector comprising transforming a host cell with a vector for polypeptide in the cell, and recovering the fusion polypeptide from the cappressing a polynucleotide sequence encoding the fusion polypeptide from the cappressing a polynucleotide sequence encoding a fusion polypeptide of cappressing a polynucleotide sequence encoding a fusion polypeptide of formula (1), (II), (III), (IV), or (V) or its salts (R_1-L-R_2 (I), R_2-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, IgE receptor; reepsilountalpin, ic., antiallergic; dermatological; antiallemmatory; antiasthmatic; 88; IgE binding domain; systemic allergy; IgE-receptor-mediated disorder; atopic dermatitis; atopic asthma; chronic urticaria; primer.
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ABS52976 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                               New fusion IgE-binding polypeptide, useful for the prevention an treatment of systemic allergy and/or other IgE-receptor-mediated disorders such as atopic dermatitis, atopic asthma and chronic u
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                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 7; 49pp; English.
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nilarity 85.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       T cell receptor; TCR; receptor; variable alpha peptide; AV peptide; TCRV; T cell variable gene; T cell regulatory activity; autoimmune disease; multiple aclerosis; human; reverse transcriptase; RT-PCR; primer; as.
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                                                                                                                                                                                                                                                                  Vandenbark AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T cell receptor (TCR) variable alpha (AV)
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                                                                                                                                                                                                                                                                                                                        12-MAY-2000; 2000US-0203984P
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Pred. No. 1
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The invention relates to a method for identifying a T cell receptor variable (TCRV) gene expressed by target T cells in an individual, comprising determining expression of TCRV genes by activated T cells from the individual and determining regulatory activity elicited in response to TCRV peptides from the individual. A preferentially expressed TCRV gene whose TCRV peptide elicits low T cell regulatory activity is identified as a variable gene expressed by target T cells. The method is used to identify disease-associated T cells in an individual so that individualised therapies can be designed to prevent or treat the disease particularly an autoimmune disease, especially multiple sclerosis. This sequence represents a reverse transcriptase PCR (RT-PCR) primer used in

T cells from

WPI; 2002-697882/75.

Identifying a T cell receptor variable gene expressed by target T cells in an individual is useful to identify disease-associated T cells for design of individualized therapies, particularly for autoimmune disease

Page 11;

English

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RESULT 2632
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Best Local :
                                                                                               The invention provides new human C2H2-type zinc finger proteins, MFQ-110. The MFQ-110 polypeptides can be expressed by standard recombinant methodology. The MFQ-110 polypeptides and polynucleotides can be used in diagnostic assays for detection of abnormally decreased or increased levels of polypeptide or mRNA expression. This may be used for diagnosing or determining susceptibility of a subject to disease that include developmental disorders, neurodegenerative disease, brain stroke, psychiatric disorders such as schizophrenia, cardiac and vascular disease, angiogenesis and cancer especially lymphomas. The polypeptides may be used to identify membrane bound or soluble receptors and may be used to identify membrane bound or soluble receptors and may be used to identify agonists and antagonists which compete with receptor binding. The polynucleotides may be used to identify membrane brained gene, for chromosome localization studies and tissue expression studies. Sequences AMI67914-17 represent
                                                                                                                                                                                                                                                                                                                                                              Identification of a new human C2H2-type finger protein, MFQ-110, which may be useful in the treatment and diagnosis of disease such as developmental disorders, neurodegenerative disease, vascular disease and
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Rosell Vives E;
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                                                                                                                                                                                                                                                                                                                        Page 62; 63pp;
                                                                                        the
                                                         BP; 4 A;
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA cloning primer
                                                                                     PCR amplification and cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Messeguer Peypoch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
             0.2%;
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                                                         4 C; 5 G; 9 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                       English.
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               Score 15.2;
Pred. No. 1.
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Pred. No. 1.9e+03
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 Mismatches
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               .9e+03
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                           Length
                                                                                      human MFQ-110
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Gaps
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RESULT 2633
ABK55618/c
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27-MAR-2001;
24-AUG-2001;
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25-AUG-2000;
25-AUG-2000;
30-AUG-2000;
30-AUG-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
22-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pancreatitis; musculoskeletal disorder; Parkinson's disease; Huntington's disease; behavioural disorder; pain; obesity; wound hearneurodegenerative disorder; neuropsychiatric disorder; hypertension; growth disorder; reproductive disorder; lung disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diabetes; cell signal processing; metabolic pathway modulation; inflammation; autoimmune disorder; scleroderma; transplantation; systemic lupus crythematosus; haemophilia; Alzheimer's, disease; graft versus host disease; Lesch-Nyhan syndrome; periodontitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human NOV3a
                                                                                                                                                                                                                  Gerlach
                                                                                                                                                                                                                                                                                  31-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                         27-AUG-2001; 2001WO-US026518
                                                                                                                                                                                                                                                                                                                                                                                                             28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                               WO200216600-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ss; primer; NOVX; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK55618;
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2000US-0228324P
2000US-0228997P
2000US-0229185P
2000US-0229780P
2000US-0229848P
2000US-0229848P
2000US-026337P
2001US-026337P
2001US-026337P
2001US-0265518P
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2001US-00939398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cardiomyopathy; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                wound healing
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allergy;

Spytek KA, Tchernev V Ĭ, Macdougall J Zerhusen BD, Padigaru Z, , Smithson G, Stone Rastelli L, Verney C Taupier RJ; CAM, Ellerman K; Patturajan 3

WPI; 2002-292064/33

musculoskeletal disorders. polypeptides, termed NOVX, useful for treati disorders, hemophilia, Lesch-Nyhan syndrome, New isolated cytoplasmic, nuclear, membrane bound and secreted, useful for treating inflammation, asch-Nyhan syndrome, pancreatitis, autoimmune

Example 2; Page 207; 245pp; English.

or secreted polypeptide, designated NoVX (actually NOV1, 2a, 2b, 3a, 3b, 4, 5a, 5b, 5c, 5d, 5e, 5f, 5g, 5h, 5i, 6, 7 and 8), a variant of NOVX, a mature form, or a variant of the mature form of NOVX. Also included are a polynucleotide encoding NOVX (or its complement), a vector comprising the polynucleotide, a cell comprising the vector, an anti-NOVX antibody, determining the presence of NOVX in a sample using the antibody determining the presence of NOVX in a sample using the antibody which binds to NOVX polynucleotide in a sample using the presence of NOVX in a sample using the probe which binds to NOVX polynucleotide, identifying a an agent which binds to NOVX (including modulators of NOVX). NOVX, the polynucleotide and the The invention relates for diagnosing, to an isolated cytoplasmic, treating or preventing nuclear, membrane bound

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8888888888888888888888888888888888888
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Best Local
 07-SEP-2001;
10-SEP-2001;
17-SEP-2001;
17-SEP-2001;
17-SEP-2001;
17-SEP-2001;
                                                                                                                                                                                                                                                                                         metabolism-related disease; obesity; central nervous system disorder; Alzheimer's disease; Parkinson's disease; epilepsy; multiple sclerosis; schizophrenia; depression; autoimmune disorder; inflammatory disorder; psoriasis; allergy; lupus erythematosus; asthma; cancer; psoriasis; allergy; lupus erythematosus; asthma; cancer; inflammatory bowel disease; rheumatoid arthritis; osteoarthritis; colon cancer; lung cancer; liver cancer; breast cancer; ovarian cancer; prostate cancer; brain cancer; melanoma; liver disease; lung disease; emphysema; obstructive pulmonary disease; haemophilia; stroke; infection; PCR; nrimer, an
                                                                                                                                                                                                                                                                                                                                                                                                                              Human; NOV; gene therapy; endocrine related disease; diabetes; metabolism-related disease; obesity; central nervous system di
                                                                                                                                                                                20-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
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nilarity 85.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein associated
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   2001US-031812OP.
2001US-031813OP.
2001US-031843OP.
2001US-0322636P.
2001US-0322781P.
2001US-0322816P.
2001US-0322817P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                             2002WO-US028539
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer #19.
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No. 1
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RESULT 2635
ACF03609
ID ACF0360
XX
ACF0360
XX
IS-SEPXX
IS-SEPXX
Human N
XX
Human;
KW Human;
KW anotial;
KW anotial;
KW anotial;
KW neurop;

15-SEP-2003

(first entry)

8AON

forward

PCR

primer SEQ

H

NO:179

ACF03609 standard; DNA;

22

Human; NOVX; cytostatic; cardiant; antiinflammatory; immunosuppressive; antiallergic; haemostatic; anti-HTV; antidiabetic; antiarteriosclerotic; ancoectic; antiasthmatic; nephrotropic; antiasthritic; hepatotropic; neuroprotective; nootropic; antibacterial; virucide; antiparasitic;

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22

3921

CTCTTGGCTTCTTTCTCCC 3940

Query Match Best Local S Matches 17

l Similarity 17; Conserv

Conservative

0

Mismatches

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Gaps

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Score 15.2; DB 1 Pred. No. 1.9e+03

Length Indels

Sequence 22

BP;

13 ₽

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0 Other; 1;

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The invention describes a new isolated polypeptide (NOVX). The NOVX cc polypeptide, nucleic acid and antibody are useful as therapeutics, associated with a human disease, which includes a pathology associated cc with NOVX polypeptide. The DNA encoding the protein is useful in gene therapy for treating the disease cor condition. In particular, the NOVX cc polypeptide or polynucleotide is useful for treating endocrine/ cc metabolism-related diseases (e.g. obesity or diabetes), central nervous system disorders (e.g. Alzheimer's disease, Parkinson's disease, ce pilepsy, multiple sclerosis, schizophrenia or depression), autoimune cc and inflammatory disorders (e.g. psoriasis, allergy, lupus erythematosus, asthma, inflammatory bowel disease, rheumatoid arthritis or osteoarthritis), cencers (e.g. colon, lung, liver, breast, ovarian, cc prostate or brain cancers, or melanoma), liver diseases (e.g. liver cirrhosis), lung diseases (emphysema or obstructive pulmonary disease), these are also useful in developing powerful assay system for functional canalysis of various human disorders, as well as in disgnostic capplications, and for monitoring the effects of drugs during clinical trials. This sequence represents a primer used to isolate DNA encoding cover luman NOV proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-SEP-2001; 2001US-0323519P.
20-SEP-2001; 2001US-0325631P.
20-SEP-2001; 2001US-0323636P.
25-SEP-2001; 2001US-0324969P.
25-SEP-2001; 2001US-0324990P.
26-SEP-2001; 2001US-0324990P.
17-APR-2002; 2002US-0373312P.
06-SEP-2002; 2002US-00236177.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rieger
Lepley
                                                                                                                                                                                                                                                                                                                                                                                                                                                         New cytoplasmic, nuclear membrane bound or secreted polypeptides (NOVX) and polynucleotides, useful in gene therapy, e.g. for treating or preventing obesity, multiple sclerosis, allergy, cancers, hemophilia, stroke or infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spytek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets
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                                                                                                                                                                                                                                                                                                                                                                                                                              Example
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Vernet CAM,
A, Leach MD,
Taupier RJ,
Edinger SR,
                                                                                                                                                                                                                                                                                                                                                                                                                            Page 494;
                                                                                                                                                                                                                                                                                                                                                                                                                            586pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
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, Ellerman K, Bergh,
Catterton E, Kekud,
Shenoy SG, Liu X,
Burgess CE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li L, Ande
Berghs C,
Kekuda R,
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Ji W, Miller CE;
igaru M, Alsobrook
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alsobrook JP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhong M;
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Matches
                                                                             Query Match
Best Local :
                                                                                                                                                                                                                                                                                   antiarteriosclerotic, anorectic, antiasthmatic, nephrotropic, virucide, antiarthritic, hepatotropic, neuroprotective, antibacterial, relaxant, antiparasitic, anticonvulsant, hypotensive, vasotropic, antiparkinsonian, vulnerary, angiogenic and antiangiogenic activities, and can be used in gene therapy and vaccines. The NOVX polypeptides and their antibodies can be used to determine the presence or absence of [1] in a sample. The NOVX polypeptides, polynucleotides encoding them, and antibodies against them, are useful in manufacturing a medicament for treating or preventing a syndrome associated with a NOVX-associated disorder such as hypertension, cardiomyopathy, atherosclerosis, cancer, diabetes, asthma, inflammation, autoimmune disorders, allergies, blood disorders, obsesity, acquired immunodeficiency syndrome (AIDS), immunoglobulin (Ig)A nephropathy, infections (e.g. bacterial, viral, parasitic), stroke, muscular infections (e.g. bacterial, viral, parasitic), stroke, muscular
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Guo X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  relaxant; anticonvulsant; hypotensive; vasotropic; antiparkinsonian; vulnerary; angiogenic; antiangiogenic; gene therapy; vaccine; cancer; cardiomyopathy; atherosclerosis; hypertension; diabetes; inflammation; autoimmune disorder; allergy; blood disorder; AllS; obesity; asthma; acquired immunodeficiency syndrome; nephropathy; cirrhosis; arthritis; Alzheimer's disease; Parkinson's disease; goitre; infection; stroke; muscular dystrophy; epilepsy; wasting disorder; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACF03547 to ACF03570 encode the human NOVX proteins (I) given in ABR: to ABR57435. (I) have cytostatic, cardiant, antiinflammatory, nootrog immunosuppressive, antiallergic, haemostatic, anti-HIV, antidiabetic, antiarteriosclerotic, anorectic, antiasthmatic, nephrotropic, virucic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New NOVX polypeptide useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis or diabetes, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 260;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gunther E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-NOV-2000;
07-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-NOV-2001;
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                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chromosome mapping, tissue typing
7413
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                                                                               Similarity
CAGCAGCAGCAGCAGCA 7432
                                                                                                                                                            22
                                                                                                                                                                                                                                    epilepsy, and other wasting disorders associated with chronic ACF03571 to ACF03644 represent PCR primers and probes for NOVX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rnandes ER,
Furtak K,
                                                                                                                                                       BP;
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Ellerman K, Alsobrook JP, Lepley DM
Edinger SR, Gangolli EA, Gorman L,
nandes ER, Vernet CAM, Tchernev VT,
Furtak K, Baumgartner JC, Colman SD;
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2000US-02465317P.
2000US-0246562P.
2000US-0246871P.
2001US-0264438P.
2001US-0264429P.
2001US-02644799P.
                                                    Conservative
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Pred. No. 1
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                                                                          1.9e+03;
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                                                                                                        DB 1;
                                                                                                                                                            Other;
                                                                                                   Length 22;
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Taupier RJ, I
Casman SJ, Sh
                                                    Indels
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Shenoy
                                                    Gaps
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CAGCAGCAGCCGCATCATCA 21

16-OCT-2003 ACF57219;

(first

entry)

0

Human LAMB3

forward

PCR primer SEQ ID NO:19

Human; mouse; skin structure; skin; laminin 5 chain gene; LAMA3; LAMB3;

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ACC58206/c

ID (ACC58206 standard; DNA; 22 BF XX

AC ACC58206;

XX

ACC58206;

XX

PCR primer used in human FCGR XX

PCR primer used in human FCGR XX

Human; FCGR2A; FcgammaRIIa; r KW

Homo sapiens.

XX

HOMO sapiens.

XX

PN

WO2003035904-A2.

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PT

11-OCT-2002; 2002WO-EP011397.

XX

19-OCT-2001; 201EP-00402718.

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(UYLI-) UNIV LILLE CENT HOSPI

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(UYLI-) UNIV LILLE CENT HOSPI

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(INNA-) INNATE PHARMA.

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(UYLI-) UNIV LILLE CENT HOSPI

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(INNA-) INNATE PHARMA.

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WPI; 2003-482053/45.

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Assessing the response of a :
 PT antibody treatment, comprise:
 PT antibody treatment, comprise:
 PT antibody treatment on non-Hocolated in the present sequence is that CC determine any correlation between the antibody treatment in non-Hocolated in the antibody treatment (seen cc site that served as an interivation between the served as an interivation between the served as an interivation between the antibody treatment. In concern the correlation with antibody treatment. In concern the 
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                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is that of a PCR primer used in human FCGR2A genotyping. Genotyping of FCGR2A-131H/R was performed in order to determine any correlation between genotype and response to therapeutic antibody treatment in non-Hodgkin's lymphoma (NHL) patients. The present (sense) primer was modified to create a BstUI site in case of R allele, and the antisense primer (see ACC58207) was modified to create a BstUI site in case of R allele, site that served as an internal control. PCR amplification of genomic DNA was performed. Amplified DNA was digested with BstUI and separated by electrophoresis, staining with ethidium bromide. The FCGR2A-131H and respectively. There was no correlation between FCGR2A-131R/H genotype and response to treatment. In contrast, FCGR3A-158V/F genotyping revealed a correlation with antibody treatment response in NHL patients
ACF57219 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Assessing the response of a subject having a tumor to a therapeutic antibody e.g., rituximab treatment, or selecting patients for therapeutic antibody treatment, comprises determining the FCGR3A158 genotype of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-OCT-2001; 2001EP-00402718
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                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                    6005 GAGGGTTTCTGGCATTTTCC 6024
                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                            17;
                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                        Conservative
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ide polymorphism; SNP; lymphoma; PCR; primer; ss
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Pred. No. 1.9e+03
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ADB81310/c
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Best Local Similarity
Matches 17; Conserv
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a method and a kit for determining the expression of mRNA or cDNA of a protein participating in the maintenance of skin structure. The method is quantitative, simple and accurate in the determination of extracellular matrix components of laminin 5 chain genes (LAMA3, LAMB3 and LAMC2, matrix metalloproteinases MMP-1, MMP-2, MMP-3 and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LAMC2; extracellular matrix component; matrix metalloproteinase; MMP-1; MMP-2; MMP-3; MMP-9; TIMP-1; TIMP-2; TIMP-3; collagen; PCR primer; 88.
                                                                                                                                                                                                                                                                    310/c
ADB81310 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A method and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-2002; 2002JP-00006797.
WPI; 2003-627612/59.
                                                                                                                                                                                                        PCR
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 6 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                           MMP-9, VII collagen, type I collagen alpha 1 chain, type I collagen alpha 2 chain, type II collagen alpha 1 chain, type IV collagen alpha 1 chain, type IV collagen alpha 1 chain, type IV collagen alpha 2 chain, TIMP-1, TIMP-2 and TIMP-3. ACF57201 to ACF57290 represent PCR primers and probes used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-NOV-2002
                                                              31-JAN-2002; 2002GB-00002206
                                                                                    30-JAN-2003; 2003WO-EP000982
                                                                                                         07-AUG-2003
                                                                                                                                                                        site
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                                                                                                                                                                                                                                                 ADB81310;
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                                         (BIOI-)
                                                                                                                                                                                                                                                                                                                                               4513
                                                                                                                                                                                  PCR;
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                                                                                                                                                                      PCR; primer; F8 scFv; antibody; anti-ligand library; directed mutagenesis.
                                                                                                                                                                                                                                                                                                                          w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         d and a kit for determination of expression of mRNA or participating in the maintenance of skin structure.
                                          BIOINVENT
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                                                                                                                                                                                                                                                                                                                          CAGGACTGGAGAAGCGTGTG 22
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                                           INI
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                                                                                                                                                                                                     amplify F8 scFv antibody to generate
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PT Making a library of anti-ligands having a cavity binding site for a pT ligand, useful in screening for a ligand.

XX Example 1; Page 16; 65pp; English.

XX Example 1; Page 16; 65pp; English.

XX This invention relates to a novel method of generating anti-ligand comprises providing combinates that have a cavity binding site. The method comprises providing comparises providing several novel anti-ligands having a cavity for binding a first ligand, and differing from the first anti-ligand in that one or more of the amino comparises and comparises are comparised by site and differing from the first anti-ligand in achieved by site are comparised by site of the cavity binding surface are replaced by those having similar but compared to the first anti-ligand is an action of the cavity binding surface are replaced by those having similar but compared to the properties. Preferably the anti-ligand is an compared to the replacements are derived from a natuarl CDR (complementarity determining region) that makes up the cavity and which binds to the ligand - an antigen. Specific anti-ligands compared to improve their binding characteristics or to better corrleate their final use. This oligonucloetide is PCR primer 9 used co amplify the continuous continuous fragment in order to generate the anti-ligand library of the livention.
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Matches Query Match Best Local 1984 TTCCTGGGAGCAGATGTTAC 2003 21 1 Similarity
17; Conserv TTCCTGGGAGCTGCTGATAC 2 Conservative 0.2%; 0 Score 15.2; Pred. No. 1 Mismatches .9e+03 DB 1; Length Indels 0 Gaps

0

Sequence

22

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6 A;

7 C;

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G; 3

T; 0 U;

0 Other;

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ADC10255
ADC10257
XX ADC
XX ADC
XX ADC
XX B9;
XX Hum
XX Hum
XX Hum
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XX Ch
12-JUN-2001;
14-JUN-2001;
15-JUN-2001;
18-JUN-2001;
19-JUN-2001;
21-JUN-2001;
22-JUN-2001;
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07-JUN-2001;
11-JUN-2001;
12-JUN-2001;
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04-JUN-2001;
06-JUN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neuroprotective; antiinflammatory; gene therapy; antisens; thyromimetic; NOVX; pathology; cancer; diabetes; obesity; endocrine disorder; CNS disorder; inflammatory disorder; chromosome mapping; tissue typing; predictive medicine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human NOVX polypeptide gene forward primer SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC10255;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC10255 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; primer;
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2001US-0297414P.

2001US-0295573P.

2001US-0297567P.

2001US-0298285P.

2001US-0298528P.

2001US-029833P.

2001US-029933P.

2001US-029939P.

2001US-02993P.

2001US-0390177P.
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                                                                                                                                                                                                                                                                                                                              2001US-0296418P.
2001US-0296575P.
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RESULT 2640
ADE76823/C
ID ADE7682
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AC ADE7682
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AC ADE7682
XX
DT 29-JAN-
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                            The invention relates to novel isolated polypeptides, mature form of the polypeptide, a sequence that is 95% identical to the polypeptide or the polypeptide comprising one or more conservative substitutions. The NOVX polypeptide is useful for treating or preventing a pathology associated with the polypeptide e.g. disorders associated with aberrant expression or activity of the polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing and predictive medicine. This sequence corresponds to a primer used to amplify and isolate the coding sequence for one of the polypeptides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-JUL-2001;
31-JUL-2001;
14-SEP-2001;
25-SEP-2001;
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10-APR-2002;
10-MAY-2002;
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22-FEB-2002;
22-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated NOVX polypeptides and nucleic acid molecules useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Agee ML, Anderson DW, Berghs C, Casman SJ, Catterton E;
Dipippo VA, Edinger SR, Eisen A, Ellerman K, Gangolli EA;
Gerlach VI, Gorman L, Guo X, Herrmann JL, Hjalt T, Ji W, Kekuda
Khramtsov NV, Li L, Liu X, Malyankar UM, Miller CE, Millet I;
Ort T, Padigaru M, Patturajan M, Pena CEA, Rastelli L, Rieger DK
Rothenberg ME, Shenoy SG, Shimkets RA, Smithson G, Spaderna SK;
Spytek KA, Stone DJ, Vernet CAM, Zhong H, Zhong M, Alsobrook JP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-FEB-2002;
21-FEB-2002;
  29-JAN-2004
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12-MAR-2002;
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27-FEB-2002;
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14-DEC-2001;
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                                                                                                                                                                                                                                          Sequence
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2001US-0301550P

2001US-0301550P

2001US-0302951P

2001US-0322297P

2001US-03374779

2001US-0341562P

2001US-0341562P

2002US-035912P

2002US-035912P

2002US-0359121P

2002US-0359121P

2002US-0359134P

2002US-0359134P
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Pred.
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lerna SK;
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ADE15462
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to a novel method of identifying water-borne organisms associated with harmful algal bloom. The method comprises fractionating an aliquot of water by size to yield several fractions, isolating but from the fractions, amplifying but by high throughput, real time PCR comprising forward and reverse genus-specific primer to yield amplified DNA amplicon, contacting the amplicon with a species-specific labelled probe, and detecting the amplicon. The method is useful for identifying water-borne organisms associated with harmful algal bloom or transported in the ballast of a ship, where the water-borne organisms are dinoflagellates selected from Pficeteria, Gymnodinium, Chattonella, Alexandrium and Aureococcus, or is a zebra mussel (Dreissena). The present sequence is that of a species-specific PCR primer which may be
                     Human; T cell receptor variable region alpha; TCRV alpha; cytokine response; CD4+ T cell; CD25+ T cell; autoimmune disease; multiple sclerosis; rheumatoid arthritis; systemic lupus erythematosus; type I diabetes; non-obese diabetes; myasthenia gravis; Grave's disease; type I typoiditis; PCR; primer; ss; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                         present
used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cary
             reverse
                                                                                                          T cell
                                                                                                                                                                                            ADE15462 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying water-borne organisms associated with harmful algal bloom comprises fractionating an aliquot of water, isolating DNA from fractions, amplifying DNA, contacting amplicon obtained with probe, ardetecting amplicon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfiesteria shumwayae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               water-borne organism; algal bloom; ship ballast; d. Pfiesteria; Gymnodinium; Chattonella; Alexandrium; zebra mussel; Dreissena; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfiesteria shumwayae-specific
                                                                                                                                                                  ADE15462;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 17; SEQ ID NO 4; 36pp; English.
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10-JUL-2002;
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                                                                                                                                                                                                                                                                                                                        17;
                                                                                                          receptor variable region alpha, RT PCR
                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                  CAGATCCTGACGTACCAGCT 2564
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                                                                                                                                                                                                                                                                  CAGATTCTGACCTATCAGCT 1
                                                                                                                                                                                                                                                                                                                                                                              BP;
                                                                                                                                                                                                                                                                                                                                                                                                          method of the invention.
                                                                                                                                                                                                                                                                                                                        Conservative
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2002US-0394654P.
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                                                                                                                                       (first
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                                                                                                                                                                                                                                                                                                                                                                              3 C; 6 G;
             PCR.
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                                                                                                                                                                                                                                                                                                                                     Score 15.2;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                              6 T;
                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                              0 U;
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                                                                                                                                                                                                                                                                                                                                                                              Other;
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                                                                                                            primer
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                                                                                                                                                                                                                                                                                                                                                 Length
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                                                                                                                                                                                                                                                                                                                        Indels
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RESULT 2642
AAT05666/c
ID AAT0566
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AC AAT0566
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DT 20-JUN-
XX
DE PCR pri
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                                                                                                                                                                                                                                                                                                                                                                The invention relates to identifying a T cell receptor (TCR) variable (V) copptide useful as a therapeutic agent in a subject with a disorder, CC comprising screening TCR V beta and/or TCR V alpha peptides to select a CC TCR V peptide that produces altered expression of a cytokine in response CC to the peptide by T cells from the subject, and determining a regulatory CC activity of CD4+CD25+ T cells isolated from the subject in response to CC the peptide. Also included are monitoring the efficacy of a TCR V peptide CC subject to the peptide and determining a T cell regulatory activity of CD4+CD25+ T isolated from the subject, where induction or regulatory activity indicates the efficacy of the peptide for treatment of the SC subject), selecting a therapy for a subject (comprising: identifying a CC RV gene expressed by target T cells in the subject by screening for CC expression of a TCR V gene by activated T cells from the subject and determining expression of a cytokine elicited in response to one or more CC subject, thereby identifying a TCR V gene expressed by target T cells for TCR V gene by T cells from the subject and clear if the subject, thereby identifying a TCR V gene expressed by target T cells or responding to the TCR V gene by target T cells)

CC subject, thereby identifying a TCR V gene expressed by target T cells)

CC nemtide is useful for identifying a T cell receptor (TCR) variable (V)

CC peptide useful as a therapeutic agent in a subject with a disorder. The
                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                      peptide is used to treat an autoimmune disease, particularly multiple sclerosis, rheumatoid arthritis, systemic lupus crythematosus, type I diabetes, non-obese diabetes, wasthenia gravis, Grave's disease, Hashimoto's thyroiditis or psoriasis. The present sequence is a TCR V alpha reverse transcriptase (RT)-PCR primer used to measure TCR gene expression levels, in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-MAY-2003; 2003US-00438729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-MAY-2000;
10-MAY-2001;
 PCR primer for nucleic acid encoding TNF microsatellite.
                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 125; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying T cell receptor variable peptides useful for treating autoimmune disease including multiple sclerosis, rheumatoid arthritis, diabetes, myasthenia gravis, Grave's disease, Hashimoto's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-864176/80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vandenbark AA;
                            20-JUN-1996
                                                                                  AAT05666 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thyroiditis and psoriasis.
                                                                                                                                                                              6195 GAGAATGGAGAGAATTTGAA 6214
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US DEPT VETERANS AFFAII
                                                                                                                                                                                                                      Similarity
                                                                                                                                                     GTGAATGGAGAGAATGTGGA 20
                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                              ВP;
                                                                                                                                                                                                           Conservative
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2001US-00853830.
                            (first entry)
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                                                                                                                                                                                                                     85.0%;
                                                                                  DNA;
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                                                                                                                                                                                                         0,
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Pred. No. 1.
                                                                                                                                                                                                                                                              G; 4 T; 0 U;
                                                                                                                                                                                                                       1.9e+03
                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                 Length 22;
                                                                                                                                                                                                            Indels
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RESULT 2643
AAT63128/C
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Best Local Sim
Matches 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT63128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Screening for Crohn's disease - by detecting nucleic acid encoding particular tumour necrosis factor micro-satellite allele(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumour necrosis factor; TNF; microsatellite; Crohn's disease; allele; inflammatory bowel disease; diagnosis; therapeutic; ulcerative colitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23
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22-SEP-1995;
22-SEP-1995;
                                                                                                               27-MAR-1997.
                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                               polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                           Promoter; glutathione S-transferase; herbicide safener;
                                                                                                                                                                                                                                                                                                                                     Glutathione S-transferase promoter PCR primer AI2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              alleles a2, b1 and c2 are absent
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                                                                                                                                                           WO9711189-A2
                                                                                                                                                                                                                                                                                                                                                                                     22-JUN-1997
                                                                                                                                                                                                                                                 ranscription factor; transgenic plant; maize; primer; PCR; olymerase chain reaction: ao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5179 CTCTGCATGTTCTCCACTTG 5198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        crcrecagerrerececare 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 8 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
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95GB-00019404
95GB-00019406
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                                                                   96WO-GB002116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.2;
Pred. No. 2
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(ZENE) ZENECA

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RESULT 2644
AAV57842/c
ID AAV5784
XX AAV5784
XX AAV5784
XX Human c
XX Human c
XX Manic-d
KW Manic-d
KW MyO-ino
XX Synthet
OS Synthet
OS Homo sa
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Best Local S
Matches 17
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AAV57842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           kDa subunit gene promoter region (see also AAT63125) suggested that element(s) conferring inducibility lay within 900 bp immediately upstream of the transcription start point (TSP). A series of fine deletion constructs were made by fusing 200 bp deleted fragments of this 900 bp region to a beta-glucuronidase marker gene. PCR primer AI2 (AAT63128) was designed to correspond to a PstI site adjacent to the TSP. It was used in combination with primers AI3, AI4, AI5 and AI6 (AAT63129-32) in PCR experiments to generate fragments of 217, 378, 570 and 760 bp, respectively. Transient transformation assays indicated that the
                                                                                                                                                                                                                                                                                                                                              Manic-depressive illness; susceptibility; genotype; chromosomal marker; polymorphic marker; chromosome :
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV57842;
                                                 New isolated IMP.18p myo-inositol monophosphatase - used to products for determining susceptibility to manic depressive as targets for preventive and therapeutic treatments.
                                                                                                                                   Detera-Wadleigh
Berrettini WH,
                                                                                                                                                                                                                                                                                                                                                                                                       Human chromosome 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                          Disclosure; Page 3; 118pp; English
                                                                                                          WPI; 1998-272247/24.
                                                                                                                                                                                                    28-OCT-1996;
                                                                                                                                                                                                                              28-OCT-1997;
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                                                                                                                                                                                                                                                                                 WO9818963-A1
                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                 18-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preliminary deletion analysis of the maize
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                                                                                                                                                                          DEPT HEALTH & HUMAN SERVICES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                    96US-0029278P.
                                                                                                                                                                                                                             97WO-US019381
                                                                                                                                SD, Gershon
Yoshikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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ene expression
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                                                                                                                                 ES, Badn
Sanders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                 Badner i
ders AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sheppard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    glutathione S-transferase plants, esp. with herbicide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2e+03;
                                                                                                                                 JA, Goldin
, Esterling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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                                                                                                                                                                                                                                                                                                                                               primer; ss
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                                                                  develop
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A method has

been developed for determining

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genotype associated with

PCR primers AAV64633-34 are used in PCR analysis of tumour necrosis factor (TNF) microsatellite alleles to identify TNF locus c. The specification describes a method for detecting the presence or absence of at least two TNF microsatellite alleles selected from TNFa10, TNFb4, TNFc1, TNFd3 and TNFe3 in a patient with Crohn's disease, where the presence of an allelic combination comprising at least two of the alleles indicates a clinical subtype of Crohn's disease having an inferior clinical response to anti-Th1 cytokine therapy. The methods and the products can be used for diagnosing clinical subtypes of Crohn's disease, that affect the gastrointestinal tract and produce similar symptoms, with

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RESULT 2645
AAV64634/c
ID AAV6463
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           increased susceptibility to manic-depressive (MD) illness. The method comprises determining the genotype of an affected individual with at least one polymorphic marker localised within the chromosomal region defined by and including markers D18843 and D18869 and determining the genotype associated with increased susceptibility to MD disorder. The method can be used for determining susceptibility to MD illness including bipolar disorder, genetic counselling of individuals from families affected with MD illness, and aid in the differential diagnosis of MD illness from other psychiatric pathologies. Products from the present invention can also be used to obtain modulators of IMP.18p myo- inositol monophosphatase protein activity and as targets for preventive and therapeutic treatments. The present sequence represents a PCR primer from Table 1 in the present invention (see AAV57798 to AAV57877)
                                                                                                                                                                   Diagnosis of Crohn's disease sub:types - by detecting serological genetic markers for identifying sub:types having characteristic responsiveness to anti-Th1 cytokine therapy.
                                                                                                                                             Claim 37;
                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                  Plevy SE,
                                                                                                                                                                                                                                                                            (CEDA-)
                                                                                                                                                                                                                                                                                                                                                         08-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tumour necrosis factor; TNF; microsatellite allele; Crohn's disease; clinical subtype; anti-Th1 cytokine therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV64634;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23
                                                                                                                                                                                                                                                                                                                 11-APR-1997;
12-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                   22-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer for
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                                                                                                                                                                                                                                                                            PROMETHEUS
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                                                                                                                                           Page 76; 117pp; English.
                                                                                                                                                                                                                                                 Targan SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 6 A; 3 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                        SINAI MEDICAL
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97US-00855825
                                                                                                                                                                                                                                                                                                                                                          166900Sn-0M86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amplification of tumour necrosis factor locus
                                                                                                                                                                                                                                                                           LAB
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                                                                                                                                                                                                                                                  Taylor
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RESULT 2646
AAV5546/c
ID AAV5554
XX AAV5554
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XX DO 9-MAR-
XX DO 9-MAR-
XX Erythro
XX Erythro
XX Erythro
XX Homo Ba
XX US58562
XX US58
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Best Local Similarity
Matches 17; Conserv
                                                                                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-OCT-1989;
12-OCT-1990;
08-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         characteristic responsiveness to anti-Th1 cytokine therapies such as anti-
                                                                                                                                                                                                                                                                        This sequence represents a PCR primer used in the creation of the human erythropoietin (EPO) isoforms of the invention. The isoforms have a single isoelectric point and have a specific number of sialic acids per molecule, the number being selected from 1-14, and the isoform being the product of the expression of an exogenous DNA sequence in a non-human eukaryotic host cell. The isolated EPO isoforms have a defined sialic acid content and biological activity, e.g. the relative in vivo specific activities increase stepwise from isoforms having 5 sialic acid residues to isoforms having 11 sialic acid residues. The EPOs can be used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV55546;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Strickland TW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer for creation of Human erythropoietin isoform.
                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Col 15; 26pp; English.
                                                                                                                                                                                                                                                     increasing
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                                                  1185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             isolated erythropoietin isoforms -
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  CTTCCGACAGGTTGGCCAGG 4
                                                  CTGCCTACAAGTTGGCCAGG 1204
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                                                                                                                                                                                                                                                           haematocrit levels in mammals
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                                                                                                                                                                                                      BP; 4 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                       Conservative
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90US-00598448.
92US-00942126.
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Pred. No. 26
                                                                                                                            Score 15.2; DB 1;
Pred. No. 2e+03;
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RESULT 2647
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ID AAXX00262;
XX AAXX00262;
XX AAXX00262;
XX AAXX00262;
XX TO Emicrosatellite 1
DE TNF microsatellite 1
XX TNF microsatellite 1
                                                                                                                                                                                                            RESULT 2648
AAX18542/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A method has been developed for diagnosing a medically resistant subtype of ulcerative colitis (UC) by detecting anti-Saccharomyces cerevisiae antibodies (Ab) in a patient. Alternatively, this subtype is diagnosed by detecting at least two of the tumour necrosis factor (TNF) alleles (microsatellite loci) TNF a2, b1, c2, d4 and e1, more specifically the TNFa2blc2d4e1 haplotype (A). Ab and haplotype (A) are useful as independent markers of the resistant subtype of UC. The methods identify a subset of UC patients who do not respond to treatment with anti-inflammatories and/or immunosuppressants, and who generally will eventually require colectomy. Barly identification of the resistant subtype will allow aggressive treatments to be applied before inflammation has become refractory. The present sequence represents a specifically claimed PCR primer for amplifying TNF microsatellite loci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TNF microsatellite locus; tumour necrosis factor; human; PCR primer; amplification; diagnosis; ulcerative colitis; TNFa2b1c2d4el haplotype; anti-inflammatory; anti-Saccharomyces cerevisiae antibody; colectomy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Diagnosing a resistant clinical subtype of ulcerative colitis - detecting presence of anti-Saccharomyces cerevisiae antibodies of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TNF microsatellite loci c
   Mouse IMC
                                                             05-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 25; Page 44; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          particular combination of tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-095224/08
                                                                                                                                                                                   AAX18542 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                               CTCTGCATGTTCTCCACTTG 5198
                                                                                                                                                                                                                                                                                                                                    CTCTGCAGGTTCTCCCCATG 2
carcinoma cell IMC-HA1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Targan SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 C; 8 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
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                                                                                                                                                                                   ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.2;
Pred. No. 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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      PCR
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   primer
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      QES.
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         NO:32
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SEXSERE

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RESULT 2649
AAA13773/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides gene sequences associated with cancer metastasis which are isolated from mouse IMC carcinoma cells by detection of their higher expression in IMC-HM cell lines than in IMC-LM cell lines using differential display of the mRNA in these cells. The gene sequences can be used for the screening of potential inhibitors of cancer metastasis by either: biringing into contact with the cancer metastasis associated protein (CMAP) and determining the degree of binding; or creating a transformant cell line which expresses CMAP and measuring the degree of expression of CMAP using an antibody recognising the protein, either in the presence or absence of the potential inhibitor. IMC-HM cells transformed with antisense CMAP DNA show a lowered ability to
                                                                                                                                         Maize; glutathione-S-transferase; GST; promoter; plant; tuber; potato; expression; sprouting inhibition; storage; PCR primer; ss.
                                                                                                                                                                                      Deleted GST
                                     13-SEP-1999;
                                                                                                                                                                                                               26-JUL-2000
                                                                                                                                                                                                                                                                  773/c
AAA13773 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse; carcinoma cell; IMC-HA1; cancer; metastasis; CMAP; inhibitor; cancer metastasis associated protein; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein associated with cancer metastasis and gene encoding it for screening for potential inhibitors of cancer metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9845431-A1
            25-SEP-1998;
                                                                06-APR-2000
                                                                                           WO200018930-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-080732/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Morita M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BANY ) BANYU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          musculus.
                                                                                                                                                                                                                                                                                                                                                              3222 TGGGAGGAGGGAAGGATTTT 3241
                                                                                                                                                                                                                                                                                                                                      23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sise. The present sequence represents a PCR primer used in an
from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                              23
                                                                                                                                                                                                                                                                                                                                     TGGGAGCAGGGAAGTAGTTT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arakawa H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 20; 74pp;
                                                                                                                                                                                   promoter sequence PCR
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 5
                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM CO LTD.
            98GB-00020970
                                     99WO-GB003021.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97JP-00105333.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 9 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                    23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Japanese.
                                                                                                                                                                                                                                                                                                                                                                                         <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.2;
Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                   primer
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RESULT 2650 AAA59333/c

AAA59333 standard; DNA; 23

В₽

AAA59333;

밁 S

23

4

EP1024196-A1

Fusarium oxysporum.

Regulatory activity; transcription; I-2 resistance egg plant; potato; melon; tobacco; Arabidopsis; platissue-specific; PCR primer; ss.

plant pathogen;

pathogen; fungi;

gene

PCR primer used to amplify the cDNA sequence of I-2 resistance

(first entry)

New I-2 resistance gene tissue-specific regulatory sequence useful

Haring

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Cornelissen

влс,

Meg

5

Simons

2000-516034/47

29-JAN-1999; 29-JAN-1999; 02-AUG-2000.

99EP-00400212 99EP-00400212

(KEYG-) KEYGENE NV.

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Best Loc
Matches
                                            Query Match
                                                                  Sequence 23
                                                                                        claimed deleted GST promoter sequence comprising 693 bases
                                                                                                                                                                                                                                                                                                                                  Constitutively expressing a target gene in a storage organ or stem plant comprises transfecting the plant with a gene promoter region the 27 kD subunit of glutathione-S-transferase operably linked to a
                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-293164/25
                                                                                                                                                                                                                                                                                                                                                                                                      Robertson NM,
                                                                                                                                                                                                                                                                            A method has been developed of constitutively expressing a target
                                                                                                                                                                                                                                                                                                   Example 1; Page 15;
                                                                                                                                                                                                                                                                                                                                                                                                                             (ZENE ) ZENECA LTD.
                     Local Sim hes 17;
7411 ATCAGCAGCAGCAGCAG 7430
                                  Similarity
                                                                  BP; 3 A;
                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       Paine
                                                                                                                                                                                                                                                                                                  53pp; English.
                                0.2%;
                                                                  7 C; 5 G; 8 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                       Jepson
                      0;
                                 Score 15.2;
Pred. No. 2
                      Mismatches
                                 2e+03;
                                                                    0 Other,
                                            DB 1;
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                                            Length
                       Indels
                                             23
                      0
                      Gaps
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RESULT 2651
AAC80271/c
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Best Local
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAA59333-34 were used to amplify cDNA encoding an I-2 resistance protein. The specification describes nucleotide sequences which have a regulatory activity on the transcription of the I-2 resistance gene in plant host cells. The transgenic plants, especially tomato, egg plant, potato, melon, tobacco and Arabidopsis, are capable of expressing a gene mediating resistance to a plant pathogen, such as fungi, in a tissue-specific manner. The plant is capable of preventing infection by a plant pathogen, such as fungi. Inserting the regulatory activity polynucleotide into plant cell genomes is useful for providing plants with reduced susceptibility to plant pathogens, especially for
                                                                                                                                                          Locus-specific, separate amplification of exon 2, exon 3, and/or exon 4 of human leukocyte antigen (HLA)-A, HLA-B, or HLA-C alleles using defined primer sets, useful for subtyping or typing of HLA Class I alleles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plant resistance mechanisms against plant pathogens such as fungi.
                                                         The present invention relates to a method for the locus-specific, separate amplification of exon 2, exon 3, and/or exon 4 of human leukocyte antigen (HLA)-A, HLA-B, or HLA-C alleles. The method is useful for subtyping or typing of HLA class 1 alleles. The present sequence is an amplification primer used in the method
                                                                                                                                                                                                                                                                                         09-APR-1999;
11-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                      19-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                               WO200061795-A2
                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                           HLA-A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reverse primer #99 used for amplification of HLA-A exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC80271;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protecting plants in cultivation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                        De Canck
                                     Sequence
                                                                                                                                       Claim
                                                                                                                                                                                                               WPI; 2000-647426/62
                                                                                                                                                                                                                                                                                                                             05-APR-2000; 2000WO-EP002998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
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                                     B₽;
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                                                                                                                                                                                                                                        Rombout
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
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99US-0138614P.
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                                       σ
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                                     Ą
                                                                                                                                     128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
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 0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          typing; primer;
                                     9 C; 4 G; 3 T; 0 U; 1 Other;
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Pred.
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Pred. No. 26
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 15.2; DB 1;
No. 2e+03;
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RESULT 2653
AAF76321/c
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AAH19012/c
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                                                                                                                                                                                Matches
                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                               Nucleic acids encoding uncoupling protein 3 (mitochondrial, proton carrier) (UCP3) proteins comprising single nucleotide polymorphisms, useful for the design of drugs for treating obesity.
                                                                                                                                                                                                                                                       The present invention relates to the human uncoupling protein 3 (mitochondrial, proton carrier) (UCP3) gene and polymorphisms. The polymorphisms are associated with obesity, especially diabetes mellitus associated obesity. They polymorphisms may be identified and analysed to determine whether an individual is susceptible to obesity and may be used as the basis for targeted design of drugs to treat obesity. The present sequence was used in the identification and amplification of UCP3
                                                                                                                                                                                                                                                                                                                                                                                                                              Chew A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH19012 standard;
                                                                                                                                                                                                                       Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UCP3; uncoupling protein 3; polymorphism; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Forward primer used to amplify UCP3 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH19012;
                  Human TNFc microsatellite marker reverse
                                         65-JUN-2001
                                                             AAF76321;
                                                                                                                                                                                                                                             polymorphisms
                                                                                                                                                                                                                                                                                                                                           Example 1; Page 33; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                           WPI, 2001-218562/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-MAR-2001.
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                                                                                  AAF76321
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (STEP/)
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                                                                                                                                                                                         Similarity
                                                                                  standard;
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                                        entry!
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Pred. No. 2
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                                                                                                                                                                                Mismatches
                                                                                                                                                                                          2e+03;
                    PCR primer.
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                                                                                                                                                                                                                          0 Other,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diabetes mellitus;
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Autoimmune disease; diagnosis; susceptibility; 2-2-4 haplotype;

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RESULT 2654
AAS12665
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC The invention relates to a method of diagnosing or predicting
CC susceptibility to an autoimmune disease in an individual. The method
CC comprises determining the presence or absence in the individual of a
CC disease-associated haplotype, i.e., the major histocompatibility complex
CC (MHC) class III 2-2-4 haplotype, or an allele associated with the MHC
CC class III 2-2-4 haplotype, where the presence of the 2-2-4 haplotype is
CC diagnostic of or predictive of a susceptibility to autoimmune disease.
CC The 2-2-4 haplotype refers to the Notch 4, HSP70-HOM and D6S273 loci
CC within the MHC class III region, where Notch 4 is present as allele 2,
CC HSP70-HOM is present as allele 2, and D6S273 is present as allele 2,
CC STP70-HOM is present as allele 2,
CC susceptibility to autoimmune diseases, including inflammatory bowel
CC susceptibility to autoimmune diseases, including inflammatory bowel
CC disease, rheumatoid arthritis, diabetes mellitus type I, and particularly
CC crohn's disease in an individual, especially an individual of of
CC predicting a variety of subtypes of Crohn's disease responsive to CA2
CC antibody therapy and other forms of anti-Th1 cytokine therapy; and
CC of the invention is non-invasive and is convenient mich these and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing susceptibility to autoimmune disease, especially Crohn's disease in individual of Ashkenazi Jewish ethnicity, by detecting three marker haplotype within the major histocompatibility complex class III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                major histocompatibility complex class III, MHC; Notch 4 allele 2; HSP70-HOM allele 2; heat shock protein 70; D6S273 allele 4; human; inflammatory bowel disease; rheumatoid architits; Crohn's disease; type I diabetes mellitus; Ashkenazi Jewish ethnicity; genotyping;
                                                                                          17-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of the invention is non-invasive, and is convenient, quick, easy and reliable. Sequences AAF76320-AAF76321 represent PCR primers used to amplify the human TNFc microsatellite marker used for genotyping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      type I diabetes mellitus; Ashkenazi Jewish ethnicity; genotyping; microsatellite marker; tumour necrosis factor; TNFc; PCR primer; ss
                             Human alpha4 integrin RT-PCR primer
                                                                                                                                                                                                                  AAS12665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-244827/25
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                                                                                                                                                                                                                                                                                                                                                                                                                                        5179
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          II; Fig 1B; 56pp;
                                                                                                                                                                                                                                                                                                                                                                                                           CTCTGCATGTTCTCCACTTG 5198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                              CTCTGCAGGTTCTCCCCATG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with and without Crohn's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                          (first entry)
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 T; 0 U; 0 Other;
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WO200164207-A2

Homo sapiens.

rejection; family; ss.

vascular cell surface adhesion molecule; CAM; immunos selectin; cadherin; autoimmune disease; inflammatory

multidrug resistance; MDR; extracellular

Bisphosphonate compound; chemotherapeutic;

radiation treatment; cancer; immunoglobulin;

integrin;

Primer used

for human alpha4 specific reactions (downstream).

16-NOV-2001

(first entry)

дан99915;

AAH99915 standard; DNA;

23

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RESULT 2655

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                                                                                                                         The invention relates to inhibiting cell adhesion mediated drug resistance (CAM-DR) in a patient involving administering to the patient, a peptide (comprising D-amino acids and having a sequence of KMVIYWKAG or its variant) that inhibits CAM-DR. The peptide is useful for enhancing efficacy of chemotherapy or radiation therapy in treating cancer (myeloma, preferably multiple myeloma) in a patient. The method improves the efficacy of chemotherapy and/or radiation therapy of cancer cells such as multiple myeloma without any disadvantages experienced during practice of conventional methods. The present sequence is an RT-PCR (reverse transcriptase PCR) primer used to amplify alpha4 integrin RNA semi-quantitatively in order to check transcription of this gene in a C cell line, KSc2/VLA-4, during an experiment where the cells (adhered to microtitre plates via interaction between the alpha4 integrin and the FN c receptor) are challenged with acute cytotoxic drugs
                                                            Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; alpha4 integrin; ss; PCR primer; cancer; multiple myeloma; cell adhesion mediated drug resistance; CAM-DR; cytostatic.
                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            뮸.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Inhibiting cell adhesion mediated drug resistance (CAM-DR) useful for enhancing efficacy of chemo- and/or radiation therapy in treating cancer such as multiple myeloma, by administering a peptide that inhibits CAM-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                        5729 CIGGCTTCCTTTCCCTTTTC 5748
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                                                                                                                                                                                                                                                                                                                                                                                          2; Page 33; 78pp; English.
                                                                             Similarity
CIGGCTICTITICCACTITC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Damiano
                                                                                                                        BP;
                                                             Conservative
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Pred. No. 2e+03;
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RESULT 2656
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ID AAC8512
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XX O8-MAY-
DT O8-MAY-
DE R. anat
XX OmpA; C
KW Septice
XX Septice
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X
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to the use of bisphosphonate compounds for inhibiting cell adhesion mediated drug resistance and enhancing efficacy of chemotherapy and/or radiation therapy in the treatment of cancer by inhibiting integrin-mediated cell adhesion. Cell adhesion is required by many normal processes but in some circumstances is undesirable, being involved in many pathologies. Cancer cell interaction with the extracellular matrix prevents apoptosis and can result in cell adhesion mediated drug resistance. This nucleotide sequence represents a primer used for human alpha4 integrin subunit specific reactions
                                                                New OmpA gene of Riemerers
for diagnosing septicemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAY-2001
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                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OmpA; outer membrane protein; avian; immunization; poultry; vaccine; septicemia anserum exsudativa; antibacterial; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC85127;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       radiation treatments.
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17; Conser
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                                                                                                                                                                                                                                                           INST
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                                                                                                                                                                                                          Sumathi
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                          Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 2 A; 9 C; 2 G; 10 T;
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                                                                                                                                                                                                                                                                MOLECULAR AGROBIOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                             of Riemerella anatipestifer for production of vaccines septicemia anserum exsudativa of avian species.
                                                                                                                                                                                                                                                                                                                   99WO-SG000075
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               compounds for inhibiting cell adhesion mediated nhancing efficacy of chemotherapeutic and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene amplifying primer 4.
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                          English
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RESULT 2657
ABS73976/c
ID ABS739776/c
ID ABS7397
XX ABS7397
XX ABS7397
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-NOV-1992;
22-NOV-1993;
06-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a Riemerella anatipestifer outer membrane protein OmpA. The OmpA protein can be expressed by standard recombinant methodology. An antibody (Ab) specific to the OmpA polypeptide is useful for diagnosing an infection by R.anatipestifer in an avian species. The OmpA gene and protein are useful for the preparation of vaccines and serodetective diagnostic assays. A vaccine composition comprising the OmpA gene, protein or Ab is useful for effective immunization of poultry against R. anatipestifer infection, especially septicemia anserum exsudativa. Sequences AAC85124-139 represent PCR primers used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Interleukin-3; IL-3; ds; haematopoietic cell; haematopoietic disorder; acute myelogenous leukaemia; AML; bone marrow transplant; neutropaenia; thrombocytopaenia; aplastic anaemia; Chediak-Higashi syndrome; systemic lupus erythematosus; leukaemia; myelodysplastic syndrome; myelofibrosis; viral infection; microbial infection; parasitic infection; myelofibrosis; viral infection; microbial infection; parasitic infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Bauer SC,
Klein BK,
                                                                                                                                                                                                                                                                                 Ex vivo expansion of stem cells e.g. hematopoietic stem cells treating hematopoietic disorders, comprises culturing the cell having human interleukin-3 mutant polypeptide and harvesting of the control of the cell having human interleukin-3 mutant polypeptide and harvesting of the cells are cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABS73976
                                                                                                                                                                                                           Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-711523/77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               stem cell;
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immune deficiency; immune disorder; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the R.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Mckearn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Abrams MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B₽;
                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-00981044.
93WO-US011197.
95US-00411795.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Braford-Goldberg
Olins PO, Paik
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microbial infection; parasitic infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                               SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                            Caparon MH, Thomas JW;
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The invention relates to ex vivo expansion of stem cells, comprises culturing stem cells with a growth medium comprising a human interleub 3 (II-3) mutant polypeptide or a polypeptide comprising an N-terminal methionine residue, alanine residue or methionine-alanine di-peptide preceding the II-3 sequence, and harvesting the cultured stem cells. I include are enhancing the efficiency of the transduction of cultured in the cultured of the transduction of cultured in the cultured of the transduction of cultured in the cultured of the transduction of the transduction of the transduction of cultured of the transduction of t

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local
New viral DNA construct having one or more tumor specific transcription factor binding sites controlling expression of the E1A open reading frame, useful in the treatment of malignant and metastatic tumors of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hematopoietic cells avoids undesirable side effects caused by treatment with current drugs. The present sequence is an IL-3-associated DNA sequence which is included in the sequence listing but is not referred to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCF; tumour specific transcription factor; neoplasm; ds; tumour; cancer; malignant tumour; metastatic tumour; liver tumour; colorectal cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Double TCF binding site promoter replacement sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                13-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-JUL-2002;
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                                                                                                                                                                  2003-221752/21.
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                                                                                                                                                                                                                                      Fuerer C,
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The invention relates to a new viral adenovirus capable of replication in

DNA construct encoding for an a human or animal tumour cell,

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   comprises one or more transcription factor (TF) binding sites operatively positioned together with the EIA open reading frame to promote expression of EIA proteins in the presence of the TF. The viral construct and/or virus is useful in the manufacture of a medicament in the therapy of patients having neoplasm, where they are capable of causing death to tumour cell. They are useful in treating malignant and metastratic tumours derived from colon cells particularly liver tumours also for treating colorectal cancers and melanomas. The present sequence represents the double tumour specific transcription factor, TCF, binding site promoter replacement sequence used to insert TCF binding sites into the human adenovirus type 5 DNA sequence
                                      testing on a pooled sample, comprising employing an INVADER detection assay. The method is useful for detecting target nucleic acid sequences in pooled samples without prior amplification of the target. The method is particularly useful for detecting an allele frequency of a polymorphism, detecting a rare mutation, or testing large numbers of blood or plasma donations to eliminate units having pathogenic (e.g. viral) contamination. This polymucleotide sequence represents a probe
                                                                                                                                                                                                                                                         Use
                                                                                                                                                The invention relates to
                                                                                                                                                                               Disclosure;
                                                                                                                                                                                                           contaminated
                                                                                                                                                                                                                           or testing
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02-OCT-2001; 2001US-0326549P.
09-MAY-2002; 2002US-00326549.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pooled sample; rare mutation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABT17614;
                                                                                                                                                                                                                                                                                                                    Fors L,
                                                                                                                                                                                                                                                                                                                                                  (THIR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                             09-MAY-2002; 2002WO-US014765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-NOV-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200290572-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Invader detection assay related probe SEQ ID No 114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABT17614 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-APR-2003
                                                                                                                                                                                                                                                         of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1536 CAATGGGATTGAGATCAAAG 1555
                                                                                                                                                                                                                        of an INVADER detection assay for testing nucleic acids in pooled les without prior amplification, e.g. for detecting rare mutations, esting large numbers of blood or plasma donations to eliminate
                                                                                                                                                                                                                                                                                        2003-120555/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   w
                                                                                                                                                                                                                                                                                                                                                  THIRD WAVE TECHNOLOGIES
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                                                                                                                                                                             Fig 9; 77pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 10 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                             unics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INVADER detection assay; allele freblood; plasma donation; pathogenic
                                                                                                                                                                                                                                                                                                                    Brow MAD,
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85.0%;
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                                                                                                                                                a novel method for performing nucleic
                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6 G,
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                                                                                                                                                                                                                                                                                                                    De Arruda Indig M,
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nic contamination; probe;
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RESULT 2660
ABZ57960
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                                                                                                                                      The present sequence is a forward primer for a polynucleotide encoding a convel human G-protein coupled receptor (GPCR) that has been characterised as a respiratory chemokine receptor. RT-PCR was used to determine expression levels of the GPCR in different tissues. The receptor was compressed in respiratory tissues and tissues related to expressed in respiratory tissues and tissues related to monocyte/macrophage migration/activation, airway remodeling, airway compressed in regulation of mucocilliary clearance, regulation of mucus inflammation, modulation of neutrophil, T-cell and eosinophil migration and regulation of epithelial cell or mast cell cand/or activation, and regulation of epithelial cell or mast cell cardivation. GPCR polypeptides (see ABP58452-53) and polynucleotides (see AB257956-57) of the invention may be useful in treatment of asthma, characteristic control constructive pulmonary disease, emphysema, chronic bronchitis, in cuter respiratory distress syndrome, cough and acute bronchitis, in cuter respiratory distress syndrome, cough and acute bronchitis, in
                                                      Query Match
Best Local S
Matches 17
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide, useful for diagnosing obstructive pulmonary disease, emphysen respiratory distress syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G-protein coupled receptor; GPCR; receptor; chemokine; human; antiasthmatic; antiinflammatory; antitussive; vaccine; PCR; primer; ss.
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                                                                                                                  Seguence
                                                                                                                                                                                                                                                                                                                                                                               Example 7; Page 44; 44pp;
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                           CTACAAGTTGGCCAGGAACA 1208
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                                                        Conservative
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                                                                                                                  7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                       85.0%;
                                                                                     0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bouhelal R,
                                                                                                                                                                                                                                                                                                                                                                               English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor forward PCR primer.
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                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.2; DB Pred. No. 2e+03;
                                                                       Score 15.2;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                           agnosing or treating e.g., asthma, chronic emphysema, chronic bronchitis or acute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seuwen K,
                                                                       2e+03;
                                                                                  DB 1;
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                                                         Indels
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RESULT 2661
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                                                            least one assay selected from the group of acute myelogenous leukaemia cell proliferation, TF-1 cell proliferation, and methylcellulose assay, cand harvesting the cultured cells. The method is useful for selective exvivo expansion of stem cells. The recombinant human interleukin-3 variant or mutant proteins are useful for treating a patient having a haematopoietic disorder, such as cancer (e.g. acute myelogenous leukaemia) or certain types of B lymphoid cancers), leukopenia, neutropenia, aplastic anaemia, Chediak-Higashi's syndrome, systemic lupus erythematosus, myelodysplastic syndrome, or myelofibrosis. The cinterleukin-3 muteins are also useful as antagonists for producing antibodies used in immunoassay and immunotherapy protocols, or for stimulating bone marrow and blood cell activation and growth before infusion into patients. This sequence represents an ex vivo stem cell expansion method associated polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-NOV-1992;
22-NOV-1993;
06-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunosuppressive; antiinflammatory; interleukin agonist 3; interleukin antagonist 3; gene therapy; ex vivo expansion of stem cell; modified human interleukin-3; cell proliferation; modified human interleukin-3; cell proliferation; TF-1 cell proliferation; methylcellulose assay; haematopoietic disorder; cancer; acute myelogenous leukaemia; B lymphoid cancer; leukopenia; neutropenia; acute myelogenous leukaemia; B lymphoid cancer; leukopenia; neutropenia; aplastic anaemia; Chediak-Higashi's syndrome; systemic lupus erythematosus; myelodysplastic syndrome; myelofibrosis; bone marrow; blood cell activation; blood cell growth; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bauer SC,
Klein BK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Selective ex vivo expansion of stem cells, useful for treating a patient having hematopoietic disorder, e.g. leukemia, neutropenia or aplastic anemia, comprises using recombinant human interleukin-3 variant or mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprising separating stem cells from other cells, culturing the cells with modified human interleukin-3 polypeptide with at least 3 times greater cell proliferative activity than native human interleukin-3 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes selective ex vivo expansion of stem cells
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93WO-US011198.
95US-00411796.
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Olins P, Paik K, Polazzi J, Thomas
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Query Match Best Local Matches

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Score 15.2; Dered. No. 2e+0; Mismatches

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RESULT 2662
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leukaemia; myelodysplastic syndrome; myelofibrosis; neutropenia; thrombocytopenia; radiation; chemotherapy; bone marrow suppression; haematopoietic deficiency; azidothymidine; AZT; alkylating agent; immunosuppressive; cytostatic; haemostatic; antirheumatic; antiarthritic; osteopathic; gene therapy; cell therapy; ex vivo expansion; stem cell; human interleukin-3 mutant; hIL-3 mutant; haematopoietic disorder; aplastic anaemia; Chediak-Higashi syndrome; systemic lupus erythematosus; immunosuppressive; osteopathic; gene t chloramphenicol; rheumatoid arthritis; antianaemic; immunomodulator; dermatological; antiinflammatory; immune disorder; infection; gp

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05-JUN-2003
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04-MAR-2002; 2002US-00090182.

24-NOV-1992; 22-NOV-1993; 06-APR-1995; 09-DEC-1996; 92US-00981044. 93WO-US011197. 95US-00411795. 96US-00764114.

(CAPA/) (EAST/) (KLEI/) (BAUE/) (ABRA/) (BRAF/) BRAFORD-GOLDBERG (
CAPARON M H.
EASTON A M.
KLEIN B K.
KLEIN B K.
MCKEARN J P.
OLING P O. BAUER ABRAMS တ 3 တ

(MCKE/)

PAIK K. THOMAS J W.

Mckearn Abrams MA, JP, Braford-Goldberg Olins PO, Paik k K, Thomas Caparon MH, Ä

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2003-678181/64.

therapy, e.g. using culturing the cells Ex vivo expansion of stem interleukin-3 f stem cells (e.g. hematopoietic cells) for gene expanded stem cells for treating thrombocytopenia, in a growth medium containing a variant or mutant of by

Disclosure; SEQ ID NO 1; 242pp; English

systemic lupus erythematosus, leukaemia, myelodysplastic syndrome, myelofibrosis, neutropenia or thrombocytopenia. The method is particularly useful for ex vivo expansion of haematopoietic cells in: (a) restoring haematopoietic cells to normal amounts in those where the number of cells has been reduced due to diseases or to culturing the stem cells with a growth medium containing a human interleukin-3 (MIL-3) mutant polypeptide. The HIL-3 mutant polypeptide has a 133, 111, 133 or 111 amino acid sequence (designated HIL-3a, HIL-3b, HIL-3c and HIL-3d, respectively), given in the specification. The method is useful for ex vivo expansion of stem cells for gene therapy. The expanded stem cells are useful for treating patients with a haematopoietic disorder e.g. aplastic anaemia, Chediak Higashi syndrome, describes ex vivo expansion of stem cells comprising cases for use

The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of the SCD (stearcyl-CoA desaturase) gene by RNA interference. Also described: (1) modulating expression of SCD genes in cells, tissue explants or organisms by introduction of siNA; (3) conjugates and/or kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA. SCD inhibiting siNAs have anorectic, antidiabetic, antiarteriosclerotic, cyrostatic and virulide activities. The siNAs can be used to modulate expression of SCD virulide activities. The siNAs can be used to modulate expression of SCD virulide activities.

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29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapeutic treatments (e.g. radiation or chemotherapy); (b) preventing treating bone marrow suppression or haematopoietic deficiencies, voccur in patients treated with e.g. azidothymiddine (AZT), alkylating agents or chloramphenicol; or (c) treating rheumatoid arthritis or ot immune disorders resulting from viral, microbial or parasitic infection This sequence represents an ex vivo stem cell expansion method associpolynucleotide.
                                                                                                                             New short
diagnosis
                                                                                                                     stearoyl-CoA
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20-SEP-2002;
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11-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                               genetic
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atherosclerosis; cancer; viral infection; drug screening;
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stearoyl-CoA desaturase; RNA interference; anorectic; antidiabetion
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2002US-0363124P.

2002US-0386782P.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guanosine 5'-diphosphofucose; GDP-fucose; Guanosine 5'-diphospho-4-keto-6-deoxymannose; cardiovascular; infection; ss. /
                                                                                                                                                                                                                                            The present invention relates to producing guanosine 5'-diphosphofucose (GDP-fucose) by employing an enzyme source that is a culture broth of microorganisms. GDP-fucose is useful as synthetic substrate of complex carbohydrates that are useful e.g. for immunotherapy for protection against cardiovascular diseases, or infections by bacteria or viruses. Guanosine 5'-diphospho-4-keto-6-deoxymannose (GKDM) is useful as an intermediate for the production of GDP-fucose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Producing guanosine 5'-diphosphofucose (GDP-fucose) useful as a substrate of complex carbohydrates for immunotherapy comprises employing microorganisms that convert guanosine 5'diphospho-4-keto-6-deoxymannose
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Helicobacter pylori; alpha-1,2-fucosyltransferase; fucose-containing sugar production; Lewis antigen:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24
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                                                                                                                                                               Alpha1,2-fucosyltransferase; fucose-containing carbohydrate; virucide; antibacterial; microbial infection; anticancer; tu
                                                                                                                                                                                                                                   B alpha1, 2-fucosyltransferase coding sequence related DNA
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                                                                                         Synthetic
                                                                                                                                           primer;
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                                                                                                                                                                                                                                                                                                                                       ABA03363;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  specification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                    g carbohydrate; cytostatic;
anticancer; tumour marker;
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RESULT 2667
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method of producing a fucose-containing complex carbohydrate, involving using a culture of a transformant expressing a protein with Bacteroides-originated alphal, 2-fucosyltransferase as enzyme source, receptor complex carbohydrate and guanosine diphosphate fucose in an aqueous medium to transfer fucose to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Expression of approximatelyal,2 fucosyltransferase producing containing complex carbohydrates as preventives or remedies c microbial infections, comprises using a transformation proced
                         New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                                                                                                                                                                                                                                                                                                 diagnosis;
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                                                                                                                                                                                                                  22-MAR-2001
                                                                                                                                                                                                                                               WO200119844-A1
                                                                                                                                                                                                                                                                                                     antiinflammatory;
                                                                                                                                                                                                                                                                                                                                   Human; CD40L;
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                                                                                                                                                                                                                                                                                                                  OL; promoter; CD40 ligand promoter; rheumatoid arthritis; antiarthritic; antirheumatic; immunosuppressive:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                               tract
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Pred. No. 2
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Best Local
        The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 31-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, antirheumatic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAR74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, antirheumatic, immunosuppressive and antirinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                 New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                                                                                                                                                                                                                                                                                                     WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive;
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ARESULT 2669
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AC AAF7493
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AC 
RESULT 2670
AAF74934
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                                                                                                                                                                                                                                                                                 The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 31-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, antirheumatic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.
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17; Conserv
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Pred. No. 2.4e+03
                                                                                                                                                                                            Score 15.2;
Pred. No. 2.
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AAF74934 standard; DNA;

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WO200119844-A1

Homo sapiens

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Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.
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diagnosis; antiarthritic; antirheumatic; immunosuppressive;
antiinflammatory; inflammatory disease; autoimmune disease;
                                                       Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis;
                                                                                                                  CD40L poly-A tract sequence SEQ ID NO:17.
                                                                                                                                                                                                                                                                                       AAF74920 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 27
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17; Conserv
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Pred. No. 2.4e+03;
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RESULT 2672
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                        WPI; 2001-244776/25
                                                                                                                                                                                                                              diagnosis; antiarthritic; antirheumatic; immunosuppressive;
antiinflammatory; inflammatory disease; autoimmune disease;
                                                                                                                                                                                                                                                         Human; CD40L;
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                                                                                                   13-SEP-1999;
                                                                                                                            13-SEP-2000;
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                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                   CD40L poly-A
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                                                                          (NYRE-) NEW YORK SOC
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17; Conserv
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                                                                                                                            2000WO-US024966.
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                                                                                                                                                                                                                                                        promoter; CD40 ligand promoter; rheumatoid arthritis;
                                                                                                   99US-0153625P.
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                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                           entry)
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Pred. No. 2.
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New altered CD40L promoter for

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the study, diagnosis and treatment

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CD40L is a factor, e.g.,

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RESULT 2673
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Best Local S
Matches 17
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           The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 31-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, antinrheumatic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CD40L poly-A tract sequence
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                                                                                                                                                   3; 90pp;
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                                                                                                                                                                                             IL promoter for use in the study, diagnosis and treatment inflammatory disorders and autoimmune diseases, such as
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                                                                                                                                                   English
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , N
                                                                                                                                                                                                                                                                                                       CRIPPLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                띪
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>س</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ligand, having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       as diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (corresponding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                given in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence
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                                               Query Match
Best Local S
Matches 17
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Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                    The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                     New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-244776/25
                                                                                                                                                                                                                                                                                                                                                                                                                                   13-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-SEP-2000; 2000WO-US024966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200119844-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; CD40L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CD40L poly-A tract sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF74927
                                                                                                  Sequence
                                                                                                                                                                                                                                                                             Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                             (NYRE-) NEW YORK SOC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4020 AAAAAAGAGAGAAAACAAAA 4039
                       4020
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N
                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
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                                                                                                   28
AAAAAAAAAAAAAACAAAA 21
                     AAAAAAGAGAGAAAACAAAA 4039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAACAAAA 21
                                                                                                                                                                                                                                                                                                                                                                                    Ľ
                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0153625P
                                                                                                  22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 3 C; 0
                                                                                                  A; 3 C; 0 G; 3 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                             RELIEF RUPTURED & CRIPPLED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CD40 ligand promoter; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G; 3 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.2;
Pred. No. 2
                                                             Score 15.2,
Pred. No. 2.
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                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NO:24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.4e+03
                                                             .4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                         DB 1;
                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28
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                                                 0
                                               Gaps
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RESULT 2675
AAQ93201/c
ID AAQ9320
XX AAQ9320
XX AAQ9320
XX Enterot
XX Enterot
XX Clostri
PA (Clost)
PT New pri
밁
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                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2676
AAQ20018/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The presence of beta 1 and beta 2 toxin genes was examined by PCR in a series of type B and C Clostridium perfringens strains. For beta 2 gene amplification, primers 8319 and 8720 (AAQ93199-200) were used; primers Betatoxi and BetatoxR (AAQ93201-02) were used for the beta 1 gene. The 3 B strains examined possessed the beta 1 gene. Type C strains had either
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Enterotoxin;
Clostridium |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ93201 standard;
            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; Page 28; 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New primers for amplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-240681/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fach P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INSP )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09517521-A2
                                                                                       Synthetic.
                                                                                                                                                         deoxyribonucleic acid; major groove; ethancamino group;
aziridinylcytosine; cross-linking group; human immunodeficiency virus;
                                                                                                                                                                                                                               Cross-linking
                                                                                                                                                                                                                                                                             01-APR-1992
                                                                                                                                                                                                                                                                                                                                                                       AAQ20018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         perfringens beta 1 toxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        new beta 2 toxin food and faecal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            beta 2 gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INST PASTEUR.
CNEVA CENT NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                       standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTTTACAGAGGAAAATAGGAAACCCTCC 7044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Guillou J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTTCAAAAAAAAAATAAAAAACCTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     beta 1 toxin; food poisoning; perfringens; polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                               agent 132 binds to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93US-00172026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94WO-EP004292.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 A;
                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or the beta 1 gene, or both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ification of Clostridium perfringens toxin genes gene, used to detect and quantify C.perfringens samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 C; 6 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Popoff M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ETUD VETERINAIRES & ALIME.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.2; D
Pred. No. 2.5e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer BetatoxL.
                                                                                                                                                                                                                                  HIV target DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;; faeces; c
reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     contamination;
; primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88
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RESULT 2677

PAQ20017/c

ID AAQ20017 standard; DNA; 15 BP.

XX

AC AAQ20017;

XX

DT 01-APR-1992 (first entry)

XX

Cross-linking agent 131 binds to

XX

KW deoxyribonucleic acid; major gro

XX

KW aziridinylcytosine; cross-linkin

KW ss.

XX

OS Synthetic.

XX

FH Key
FT modified_base 2
FT /*tag= a
FT /mod_base= m5c
FT modified_base 4

/*tag= b
                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 15
                                                                                                           deoxyribonucleic acid; major groove; ethanoamino group; aziridinylcytosine; cross-linking group; human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                   This sequence is designed to bind to the HIV target sequence AGAGAGAAAAAGAG-3'and to covalently cross-link to it via the ethanocytosine (aziridenyl) group. See also AAQ20009-Q20025
                                                                                                                                                                                                                                                                                                                                                                                                                                                   New sequence-specific non-photo-activated crosslinking agents - bind to the major groove of duplex DNA and are esp. useful for treating latent infections e.g. {\rm HIV}.
                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 0 A; 5 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-007480/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAY-1990;
14-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                               Cross-linking agent 131 binds to HIV target DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 24; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAY-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-DEC-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9118997-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GILE-) GILEAD SCIE INC
                                                                                                                                                                                                                                                                                              4017 GAGAAAAAAGAGAGA 4031
                                                                                                                                                                                                                                                                         15
                                                                                                                                                                                                                                                                                                                     1 Similarity
15; Conserv
                                                                                                                                                                                                                                                                       Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Krawczyk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90US-00529346.
91US-00640654.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90US-00529346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag=
/mod_ba
                     /mod_base= m5c
                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "N4N4-ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dod.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               *tag=
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                                                                                                                                                                                                                                                                                                                                           0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     m5c
                                                                                                                                                                                                                                                                                                                   Score 15; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                    1.3e+03;
hes 0;
                                                                                                                                                                                                                                                                                                                                         DB 1; Length 15
                                                                                                                                                                                                                                                                                                                     Indels
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N4N4-
                                                                                                                                                                                                                                                                                                                     Gaps
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                               PCR; selection; primers; OPTIPRIM; breeding; genetic mapping; traits; amplification; ss.
                                                                                                                                        Microsatellite
                                                                                                                                                         25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                AAQ33752 standard; DNA; 15
                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 0 A; 5 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                         This sequence is designed to bind to the HIV target sequence AGAGAGAAAAAAGAG-3'. See also AAQ20009-Q20025
                                                                                                                                                                                                                                                                                                                                                                 New sequence-specific non-photo-activated crosslinking agents - the major groove of duplex DNA and are esp. useful for treating infections e.g. \rm HIV.
                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1992-007480/01.
                                                                                                                                                                                                                                                                                                                                                                                                                    Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAY-1990;
14-JAN-1991;
                                15-JAN-1991;
                                                                06-AUG-1992.
                                                                                WO9213102-A1
                                                                                                Bos taurus.
                                                                                                                                                                                 AAQ33752;
                                                                                                                                                                                                                                                                                                                                                  Example 4; Page 24; 42pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAY-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-DEC-1991.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9118997-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                    (GILE-) GILEAD SCIE INC.
                                                                                                                                                                                                                                                    4017 GAGAAAAAAGAGAGA 4031
                                                                                                                                                                                                                                          15 GAGAAAAAAGAGAGA 1
                                                                                                                                                                                                                                                                           15;
                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                   Krawczyk S;
                                91US-00642342
                                                                                                                                        sequence from clone TGLA162.
                                                92WO-US000340
                                                                                                                                                                                                                                                                                                                                                                                                                                                     91US-00529346.
91US-00640654.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90US-00529346
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/mod_base=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                        100.0%; **
                                                                                                                                                                                                                                                                                  0.2%; Score 15;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       m5c
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                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     m5c
                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                          DB 1; Length 15;
1.3e+03;
hes 0; Indels
                                                                                                                        cattle; parentage;
                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                    51
                                                                                                                                                                                                                                                                                                                                                                           bind to
                                                                                                                                                                                                                                                                          Gaps
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Georges M, Massey JM,

(GENM-) GENMARK.

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RESULT 2679
AAQ30250/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequence is that of a bovine microsatellite sequence obtd. by coreening a library of bovine MboI DNA fragments of between 250 and 500 cc by with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 cc clones cross-hybridised. Assuming independent distribution of cc in the bovine genome is estimated at 110, 000. The sequence information cc for ca. 230 such bovine microsatellites is summarised in the cc specification and indexed herein (see below). The sequences upstream and cc downstream of the microsatellite sequence were used to generate the crequired PCR primers for in vitro amplification of the corresp. cc microsatellite (using the program opTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic cc mapping of economic trait loci, or genes involved the determinism of ce comomically important traits esp. in cattle, to allow selective consecutive. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN cc field.)
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                    )250/c
AAQ30250 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mapping,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-284684/34
                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                         Human immunodeficiency hepatitis; malignancy;
                                                                                                                                                                                                                                                                                                                                             Oligomer HIV132 for forming triplex with HIV target duplex.
                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
07-DEC-1992
11-JUN-1992
                                                                                      modified_base
                        WO9209705-A1
                                                                                                                            modified_base
                                                                                                                                                                 modified_base
                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7415
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCAGCAGCAGCA 7429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 231; 517pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCAGCAGCAGCAGCA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bovine DNA markers - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                              _mod_
                                                 /mod_base= OTHER
/note= "OTHER= N
                                                                                                                /*tag=
                                                                                                  mod_base=
                                                                                                                                      mod_base=
                                                                                                                                                                                                                   mod_base= m5c
                                                                            *tag=
                                                                                                                                                      *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                           virus; AIDS; modified; HIV; hepatitis;
inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ВÞ
                                                                                                                                          m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                  N4 N4 ethanocytosine'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                        herpes;
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RESULT 2680
AAQ30249/c
ID AAQ3024
XX AAQ3024
XX 25-MAR-
DT 25-MAR-
DT 07-DEC-
XX Oligome
XX Human i
KW hepatit
XX hepatit
XX Synthet
XX Key
FH Key
FT modifie
FT modifie
FT modifie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   문
                                                                                                                                                                                                                                                                                                                                                                                                The synthetic oligomer is capable of forming a triplex at physiological CC pH with a purine rich target sequence by coupling into the major groove co of the duplex. The specific target sequence of this oligomer is an HIV c target duplex contg. a purine-rich region concentrated on one chain of the duplex. The oligomer, and others like it are useful in diagnosis and ctherapy of diseases characterised by specific DNA duplex targets, e.g. CHV, hepatitis, herpes, malignant tumours and inflammation. The triple chalices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. Additional modifications, such as altered inter- nucleotide linkages may also be incorporated, rendering the oligomer e.g. stable to nuclease activity. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-C 2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New oligomers contg. modified bases - doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-APR-1991;
17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 12; Page 65; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Froehler B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-SEP-
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17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-NOV-1990;
18-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-NOV-1991;
                                         modified_base
                                                                                   modified_base
                                                                                                                              Synthetic
                                                                                                                                                                       Human immunodeficiency virus; AIDS; modified; HIV; hepatitis; herpes;
                                                                                                                                                                                                  Oligomer HIV131 for forming triplex with HIV target duplex
                                                                                                                                                                                                                              25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                      AAQ30249;
                                                                                                                                                                                                                                                                                                  AAQ30249 standard; DNA; 15
                                                                                                                                                         hepatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GILE-)
                                                                                                                                                                                                                                                                                                                                                                                                   4017 GAGAAAAAAGAGAGA 4031
                                                                                                                                                                                                                                                                                                                                                                        15
                                                                                                                                                                                                                                                                                                                                                                                                                              15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                        GAGAAAAAAGAGAGA 1
                                                                                                                                                         malignancy; inflammation; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Krawczyk S,
                                                                                                                                                                                                                              (revised)
(first en
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91US-00686546.
91US-00686547.
91US-00766733.
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91US-00643382.
91US-00683420.
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  /*tag= b
/mod_base= m5c
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                                                                                                   Location/Qualifiers
                                                      mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; --
                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matteucci MD,
                                                            m5c
                                                                                                                                                                                                                                                                                                     ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                              1.3e+03;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Milligan
                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                0
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modified_base

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AAQ79185/c
ID AAQ7918
XX
AC AAQ7918
XC AAQ78
XC AAC78
XC AAQ78
XC AAQ78
XC AAQ78
XC AAQ78
XC AAC78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is an HIV target duplex contg a purine-rich region concentrated on one chain of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HIV, hepatitis, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. Additional modifications, such as altered inter- nucleotide linkages may also be incorporated, rendering the oligomer e.g. stable to nuclease activity. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JAN-1991;
08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
Nuclease resistant oligonucleotide; inhibition of gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                                           AAQ79185 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 65; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
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                                                 Nuclease resistant oligonucleotide.
                                                                                                      25-MAR-2003
21-JUN-1995
                                                                                                                                                                               AAQ79185;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-JUN-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9209705-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-217083/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GILE-)
                                                                                                                                                                                                                                                                                                                                                                                                           4017
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Krawczyk S,
                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
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91US-00683420.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mod_base= m5c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 C; 0 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       350
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.3e+03;
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RESULT 2682
AAQ79184/c
ID AAQ7918
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                                         13-OCT-1994.
                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                        Nuclease resistant oligonucleotide; inhibition 9-methyl-8-acyclo-adenosine; antisense agents;
                                                                                                                                                                                                                                                                                                                                                                                                  Nuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
21-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               184/c
AAQ79184 standard; DNA;
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                                                                                                                                                                                                         modified_base
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                                                                                                                                                                                                                                                                                                                                                                                             resistant oligonucleotide
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(first en
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                                                                                                                                    "9-methyl-acyclo-adenosine"
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                   expression;
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21-MAR-1994;

94WO-US002995

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RESULT 2683
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Best Local Similarity
Matches 15; Conserv
 15-APR-1994;
15-APR-1994;
18-MAY-1994;
06-JUL-1994;
15-AUG-1994;
16-AUG-1994;
17-AUG-1994;
                                                                                       23-FEB-1994;
29-MAR-1994;
04-APR-1994;
07-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ79182-Q79186 contain one or more 9-methyl-acyclo-adenosines, acylic nucleoside analogues which inhibit nuclease degradation. The nuclease resistant oligonucleotides can themselves be used to inhibit gene expression as antisense agents, in nucleic acid sequencing and diagnostic assays. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                   translocation; chronic myelogenous leukaemia; CML; cancer; translocation; chronic myelogenous leukaemia; CML; cancer; philadelphia chromosome; inflammation; autoimmune disease; atheroscie; myocardial infarction; stroke; restenosis; transplant rejection; sheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
                                                                                                                                                                                                                                                                                                                                               Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New acyclic nucleoside analogues - used to prepare nuclease resistant oligo-nucleotide(s) used partic. for inhibiting gene expression.
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25-MAR-1997
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                                                                                                                                                       23-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; RNA; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                hammerhead ribozyme target sequence (nt. position 2910).
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(first entry)
94US-00201109
94US-002219534
94US-0022795
94US-00224483
94US-00227958
94US-00227958
94US-00245736
94US-00291280
94US-00291932
94US-00291433
94US-00292620
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 0 T; 0 U; 0 Other;
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25-MAR-2003 25-MAR-1997

(revised) (first entry)

gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RUN; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; (NL; cancer; philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis;

Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; Human ICAM hammerhead ribozyme target sequence (nt. position 2911).

transplant rejection; rheumatoid

arthritis; psoriasis;

RESULT 2684 AAT52138

AAT52138 standard; RNA; 15

ВP

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15 DODGOODOOOOOO

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                         Query Match
Best Local S
Matches 0
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02-SEP-1994

08-SEP-1994

23-SEP-1994

23-SEP-1994

23-SEP-1994

03-CCT-1994

07-CCT-1994

11-CCT-1994

11-CCT-1994

11-CCT-1994

11-CCT-1994

11-CCT-1994

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11-CCT-1994

11-CCT-1994
                                                                                             The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the mucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in pattents with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                      Grimm
Modak
                                                                                                                                                                                                                                                                             Ribozymes having modified bases and methods for producing them in inhibiting disease related genes.
                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                 WPI; 1995-351090/45.
                                                                                                                                                                                                                                                                                                                                                                             Stinchcomb
                                                                                                                                                                                                                                                        Claim
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   4464 TTTTTTTTTTTTTT 4478
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                          Similarity 0; Conserv
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Pavco P,
                                                                                                                                                                                                                                                                                                                                          Usman
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                                                                        BP; 0 A;
                           Conservative
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94US-0030000
94US-0031039
94US-00311486
94US-00311749
94US-00316771
94US-00319492
94US-00319492
94US-00334847
94US-00337608
94US-00345516
94US-00345516
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Beigleman L,
Wincott FE,
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                           15;
                                                                        G; 0 T; 15 U; 0 Other;
                                        Pred
                                     Score 15; DB 1;
Pred. No. 1.3e+03;
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                            Mismatches
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Matulic-Adamic J, Mcswiggen JA;
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                                                   Length 15;
                            Indels
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Query Match
Best Local
  Matches
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07-OCT-1994;
11-OCT-1994;
04-NOV-1994;
10-NOV-1994;
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23-SEP-1994;
23-SEP-1994;
28-SEP-1994;
                                                              The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                                                   Ribozymes having modified bases and in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                     Modak
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06-JUL-1994;
                                              Sequence 15
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19-AUG-1994;
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16-AUG-1994;
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15-APR-1994;
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human immunodeficiency virus; acquired immune deficiency s
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           Local
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Similarity 0; Conserv
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                                                                                                                                                                                                             Page 175; 407pp; English.
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Usman N,
                                                                                                                                                                                                                                                                                                               Karpeisky
                                                                                                                                                                                                                                                                                                                           DT,
                                              BP; 0
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94US-00245736

94US-00291932

94US-00291932

94US-00293520

94US-00393520

94US-0030000

94US-00311486

94US-00311749

94US-00311749

94US-00316771

94US-00316771

94US-00337608

94US-00337608

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94US-00345516

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94US-00357577

94US-00363233
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94US-00218934.
94US-00222795.
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                                             Α,
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                                            0 C; 0 G; 0 T; 15
          0.2%;
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            Score 15;
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  Mismatches
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, Matulic-Adamic J, Mcswiggen JA;
  Sullivan SM, Sweedler D, Thompson
Woolf T;
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                        DB
                                            U; 0 Other,
          DB 1;
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4464 TTTTTTTTTTTTT 4478

RESULT 2686 AAV06038/c ID AAV0603 XX

AAV06038 standard; DNA; 15

ВP

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RESULT 2685
AAV06037
ID AAV0603
XX AAV0603
XX Oligonu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Garbesi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide-anthracycline
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08-APR-1998
                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-470805/43
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                                                                                                                         Similarity
                                                                                                                                                                                                15 BP; 10 A; 0 C; 5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 19;
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/note= "conjugated via a linker molecule
or anthracylinone"
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                                                                                                                      0.2%; Score 15; 100.0%; Pred. No.
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                                                                                                                                                                                                0 T; 0 U; 0 Other;
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                                                                                                                           1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anthracycline intercalation;
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08-APR-1998
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Arcamone F;
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                                                                                        Sequence
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                                                                                                                                                                                                                                                                      New oligo:nucleotide-anthracycline or form triple-helix complexes with DNA,
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                      4017
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                                            Conservative
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(first entry)
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                                                                                                                                                                                                                                                                      anthracyclinone conjugates - which used for targetting e.g. tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a linker molecule
                                                                  /J. 8d
                                                       1.3e+03
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                                                                  Length 15;
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RESULT 2687
                                                                                                                                                                                                                       A new method is provided for the synthesis of oligonucleotides having N3'
->P5' phosphoramidate linkages. The method comprises (a) attaching a 3'-
protected amino nucleoside to a solid support; (b) deprotecting the 3'-
amino; (c) reacting with a 3'-protected aminonucleoside-5'-
phosphoramidite monomer to form an internucleoside N3'->P5'
phosphoramidite link; (d) oxidising this link to phosphoramidate; and
coptionally repeating steps (b)-(d) until the required oligonucleotide is
completed. This method provides better yields with lower reagent
consumption than known processes, and can be operated on a large scale.
The obtained oligos, containing phosphoramidate linkages, have favourable
binding properties, nuclease resistance and solubility, and are useful as
research, diagnostic and therapeutic agents. The present sequence is an
example of an oligonucleotide in which N3'->P5' phosphoramidate linkages
have been introduced by the new method. (Updated on 25-MAR-2003 to
                                                                                Best Loc
Matches
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31-MAR-1998
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                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                            correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 28; 60pp; English.
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                                                                                                                                                                     15 A;
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/note= "these residues have N3'->P5' phosphoramidate
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Pred. No.
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thes 0;
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                                                                                    Indels
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RESULT 2688 AAV01603 ID AAV0160

AAV01603 standard; DNA; 15

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RESULT 2689
AAV07431
ID AAV0743
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AC AAV0743
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                 AAV07431;
                                                                                                                                                                                                                                                                                consumption than known processes, and can be operated on a large scale. The obtained oligos, containing phosphoramidate linkages, have favourable binding properties, nuclease resistance and solubility, and are useful as research, diagnostic and therapeutic agents. The present sequence is an example of an oligonucleotide in which N3'->P5' phosphoramidate linkages have been introduced by the new method. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthesis of N3' to P5' phosphoramidate oligo:nucleotide - by reac immobilised 3'-amino nucleotide with new amino:nucleoside 5'-phosphoramidite then oxidation, useful as research, diagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide containing phosphoramidate linkages
                                        AAV07431 standard; DNA;
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31-MAR-1998
                                                                                                                                                                                                                                         Sequence 15 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapeutic agents.
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                                                                                                                                                  4464 TTT
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                                                                                                                                                                                                                                                                      PR field.)
                                                                                                                                                                                             Similarity
                                                                                                                      TTTTTTTTTTTTT 15
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                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              THERAPEUTICS
                                                                                                                                     TTTTTTTTTT 4478
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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/note= "these residues have N3'->P5' phosphoramidate
                                                                                                                                                                                                                                       A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                          0.2%;
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                                                                                                                                                                              0;
                                                                                                                                                                                            Score 15;
Pred. No.
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1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The sequence is that of an oligonucleotide primer which was produced as CC part of an oligonucleotide peptide conjugate as an example of a release CC tag compound (RTC). These comprise a reactive group, a release group and a non-volatile mass label comprising a synthetic polymer or biopolymer CC detectable by mass spectrometry. The RTCs can be used as probes for the CC detection of TMs. They can be used for e.g. identification of gene CC sequences, identification of non-coding nucleotide sequences, detection of mutations within a gene or protein sequence, detection of metals, detection of toxins, detection of receptors on an organism or CC of metals, detections and characterisation of niteractions, enzymence substrate interactions and characterisation of ligand interactions. Multiplex applications include multiple pathogen diagnostics, multigene CC substrate interactions include multiple pathogen diagnostics, multigene CC genetic polymorphism screening, single nucleotide polymorphism (SNP) CC genotyping, clone and gene mapping, and gene expression analysis. The CR permit the ready detection of releasable mass labels by mass spectroscopy. The releasable mass labels permit the multiplexing of tens, hundreds and perhaps even thousands of different mass labels that can be considered to uniquely identify each desired target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local (
                         Capillary afinity gel electrophoresis; separation; polymer-gel; polyacrylamide; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New release tag compounds for detecting target molecule(s) - comprising reactive group, a release group and a releasable non-volatile mass label detectable by mass spectrometry.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide; peptide; conjugate; release tag compound; mass spectrometry; detection; identification; diagnosis; primer;
                                                                                                           Oligonucleotide linked to polyacrylamide.
                                                                                                                                                                 04-JUN-1998
                                                                                                                                                                                                                    AAT86675
                                                                                                                                                                                                                                                                          AAT86675 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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16-MAY-1997;
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15; Conserv
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97US-0046719P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 C; 0 G; 15 T; 0 U; 0 Other;
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Pred. No.
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC This sequence represents an oligonucleotide receptor molecule covalently CC bound to a polyacrylamide gel via a linking group. The invention relates to selective separation of electrically charged target molecules in an CC analytical mixture. It comprises capillary affinity gel electrophoresis CC using a capillary tube which is at least partly filled with a polymer CC gel. Receptors for target molecules are covalently bound to the polymer. CC An electric field of at least 50 volts/cm is applied. The capillary tube is charged with the analytical mixture. In a first separation stage, the CC is charged with the analytical mixture are bound to the receptors and the remaining components are eluted, optionally whilst splitting open. In a second stage, the elution conditions are changed, optionally in stages, co that the affinity of the traget molecules for the receptor is that the affinity of the traget molecules for the receptor is californiated and the target molecules are eluted and detected, optionally whilst splitting open. The process is useful for selective separation CC and/or determination of charged organic compounds, such as a for carbohydrares. It may be used a c for carbohydrares.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity Matches 15; Conserv
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotides, peptides or carbohydrates. It may be used, e.g. for isolation of specific proteins and DNA molecules, purification of antibodies, analysis of antisense compounds or screening for enzyme inhibitors. The process achieves higher resolution and selectivity than prior art processes, especially in the case of complex biological analytical mixtures. It has high sensitivity, even with small amounts of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                 Oligonucleotide separated by capillary affinity gel electrophoresis
                                                                                                    04-JUN-1998 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The derivatised polymers may be synthesised specifically using
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC process using capillary affinity gel electrophoresis. The invention CC process using capillary affinity gel electrophoresis. The invention CC crelates to selective separation of electrically charged target molecules CC in an analytical mixture. It comprises capillary affinity gel control of the capillary tube is charged with the analytical mixture. In a first CC separation stage, the target molecules in the mixture are bound to the capillary tube is charged molecules in the mixture are bound to the control of the capillary open. In a second stage, the elution conditions are changed, coptionally in stages, so that the affinity of the target molecules for the receptor is eliminated and the target molecules are eluted and control of c
                                                                                                                                                                                                                                                                              RESULT 2692
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Separation of electrically charged target molecules - by capillary affinity gel electrophoresis using polymer-gel to which receptors
Synthetic
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                                        Oligonucleotide; phosphoramidate; phosphoramidite; nucleoside;
                                                                                          N3-P5 phosphoramidate oligonucleotide #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard methods
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                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97WO-EP002647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15; DB 1; 
; Pred. No. 1.3e+
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                  88.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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AAX00788/c
                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                         This sequence represents an example of an oligonucleotide containing novel 3'-amino-5'-phosphoramidite nucleoside of the invention. The sequence is generated synthetically by using an amine-exchange reaction of phosphoramidites in which a deprotected 3'-amino group of an oligonucleotide chain is exchanged for the amino portion of a 5'-phosphoramidite with a protected 3' amino group. The resulting phosphoramidite internucleotide linkage is oxidised to form a stable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
misc_difference
                                                                                                                                                                            13-APR-1999
                                                                                   misc_difference
                                                                                                                                     Oligonucleotide;
                                                                                                                                                      N3-P5 phosphoramidate oligonucleotide #4.
                                                                                                                                                                                               AAX00788;
                                                                                                                                                                                                                 AAX00788
                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                  protected phosphoramidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New 3'-protected-amino-nucleoside-5'-phosphoramidite the synthesis of oligo-nucleotide(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-120007/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-FEB-1996;
14-JUN-1996;
20-DEC-1996;
                  12-JAN-1999.
                                     US5859233-A.
                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5859233-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (LYNX-) LYNX
                                                                                                                                                                                                                                                                                      4464
                                                                                                                                                                                                                                                                                                         15; Conserv
                                                                                                                                                                                                                                                                  Н
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SM,
                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                               15 BP; 0 A; 0 C; 0 G; 15 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Col 33; 34pp;
                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          THERAPEUTICS
                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nelson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-00603566.
96US-00663918.
96US-00771789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96US-00771789
                                                                                    Location/Qualifiers
1. .15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
1. .15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/note= "contains internucleotide N3-P5 phosphoramidate
/note= "contains internucleotide N3-P5 phosphoramidate
                                                  /*tag= a
/note= "contains internucleotide N3-P5 phosphoramidate
internucleotide linkages"
                                                                                                                                     phosphoramidate;
                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JS,
                                                                                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                  15
                                                                                                                                                                                                                                                                                     4478
                                                                                                                                                                                                                                                                  15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INC.
                                                                                                                                                                                                                                                                                                                                                                   linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mccurdy
                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                  Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                    phosphoramidite;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NS,
                                                                                                                                                                                                                                                                                                                                               U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hirschbein BL,
                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                 1.3e+03;
                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                           Length 15;
                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                     nucleoside;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      monomers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schultz
                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      - used
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AAX84262/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence represents an example of an oligonucleotide containing novel 3'-amino-5'-phosphoramidite nucleoside of the invention. The sequence is generated synthetically by using an amine-exchange reaction of phosphoramidites in which a deprotected 3'-amino group of an oligonucleotide chain is exchanged for the amino portion of a 5'-phosphoramidite with a protected 3' amino group. The resulting phosphoramidite internucleotide linkage is oxidised to form a stable protected phosphoramidate linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-FEB-1996;
14-JUN-1996;
                                      Protein inhibiting apoptosis, useful Alzheimer's disease.
                                                                      WPI; 1999-395181/33
                                                                                                                                                                                                        WO9931239-A1
                                                                                                                                                                                                                                                                   Nck associated protein 1; Nap1; human; apoptosis; Alzheimer's disease;
                                                                                                                                                                                                                                                                                          PCR
                                                                                                                                                                                                                                                                                                                                                      AAX84262 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 15 A; 0 C; 0 G; 0 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New
                                                                                        Sakaki Y;
                                                                                                                                          15-DEC-1997;
                                                                                                                                                               14-DEC-1998;
                                                                                                                                                                                    24-JUN-1999.
                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                         therapy; PCR primer;
                                                                                                                                                                                                                                                                                                              08-SEP-1999
                                                                                                                                                                                                                                                                                                                                  AAX84262;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 10; Col 33; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-120007/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fearon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gryaznov
                                                                                                             (KYOW ) KYOWA HAKKO KOGYO KK
(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (LYNX-) LYNX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3'-protected-amino-nucleoside-5'-phosphoramidite synthesis of oligo-nucleotide(s).
                                                                                                                                                                                                                                                                                         primer
                                                                                                                                                                                                                                                                                                                                                                                                                             4464 TITTTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                                                                                           15
                                                                                                                                                                                                                                                                                                                                                                                                                                                  15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MS,
                                                                                                                                                                                                                                                                                          for
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               THERAPEUTICS
                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nelson
                                                                                                                                                                                                                                                                                       human Nck associated protein 1 coding sequence
                                                                                                                                           97JP-00363183
                                                                                                                                                                98WO-JP005646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-00603566.
                                                                                                                                                                                                                                                           88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mccurdy
                                                                                                                                                                                                                                                                                                                                                        뫔
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SN,
                                                 ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hirschbein
                                                 the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.3e+03;
                                                 diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        monomers
                                                 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schultz
                                                treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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This sequence represents a PCR primer used

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isolate

DNA encoding

the

Example 1; Page 77; 90pp; Japanese.

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RESULT 2696
AAZ36369/c
ID AAZ3636
XX
AC AAZ3636
XX
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                                                                                                                                                                  RESULT 2695
AAX84261
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                                                                                  밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                          Matches
                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human Nck associated protein 1 (Napl) of the invention. Napl inhibits apoptosis. The protein can be used in the investigation, diagnosis and treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 0 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
          AAZ36369;
                                                                                                                                                                                   This sequence represents a PCR primer used to isolate DNA encoding the human Nck associated protein 1 (Nap1) of the invention. Nap1 inhibits apoptosis. The protein can be used in the investigation, diagnosis and treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                   14-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                        24-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                           WO9931239-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nck associated therapy; PCR pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX84261 standard; DNA;
                                                                                                                                                                 Sequence 15 BP; 5 A; 5 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         WPI; 1999-395181/33
                                                                                                                                                                                                                                                                                                                                                              15-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer for human Nck associated protein 1 coding sequence.
                                AAZ36369 standard; DNA; 15
                                                                                                                                                                                                                                                            Alzheimer's disease.
                                                                                                                                                                                                                                       Example 1; Page 77; 90pp; Japanese.
                                                                                                                                                                                                                                                                    Protein inhibiting apoptosis, useful in
                                                                                                                                                                                                                                                                                                                                (КУОМ ) КҮОМА НАККО КОĞYO KK
(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7413
                                                                                                     7413 CAGCAGCAGCAGCAG 7427
                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens.
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                                                                                                                                    Similarity
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                                                                                   CAGCAGCAGCAGCAG
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                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein 1; Napl; human; apoptosis; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                               97JP-00363183
                                                                                                                                                                                                                                                                                                                                                                                   98WO-JP005646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%; Scc.
/ 100.0%; Pr
                                                                                                                                   0.2%; Score 15;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                 ВP
                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.3e+03;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                              DB 1;
                                                                                                                                    1.3e+03;
                                                                                                                                                                                                                                                                    the diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
                                                                                                                                              Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                      and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                         0
                                                                                                                         Gaps
                                                                                                                                                                                                                                                                      of.
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                                                                                                                                                                           RESULT 2697
AAZ61854
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204-NOV-1999.

WO9955847-A2

Hepatitis C virus

Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavage; cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.

HCV 3' non core region substrate for Hammerhead ribozyme HCV.3-118

28-MAR-2000 AAZ61854;

(first entry)

AAZ61854 standard; RNA; 15

15

CAGCAGCAGCAGCAG 1

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S
                                 Query Match
Best Local Sim
Matches 15;
                                                                                                                                 The present sequence represents PCR primer used to amplify cDNA encoding a mouse cystatin-related protein, designated testatin. The protein is capable of inducing testis formation in foetal gonads. It is highly probable that the protein inhibits testis tumour growth because of structural and functional similarities with tumour inhibiting cystatins. The cystatin-related protein testatin may be useful for inducing testis formation in foetal gonads. Testatin polymucleotides are useful as a source of primers and probes, which can be used to detect the presence of testatin nucleic acid molecules in a sample. The testatin polymucleotides, polypeptides, and compositions can be used for treating genital tumours and may also be useful for creating a model for studying testic marting and may also be useful for creating a model for studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse;
foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genital tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-FEB-2000
                                                                                           Sequence 15 BP; 0 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                       Example 1; Page 15; 37pp;
                                                                                                                                                                                                                                                                                                                                                      treatment.
                                                                                                                                                                                                                                                                                                                                                                   Novel cystatin-related
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-039071/03.
                                                                                                                                                                                                                                                                                                                                                                                                                               Nordqvist K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9958565-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (KARO-) KAROLINSKA INNOVATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-NOV-1999.
                                                                                                                           cestis malformations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer used
   7413 CAGCAGCAGCAG 7427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cystatin-related protein; testatin; testis formation;
gonad; testis tumour growth; tumour inhibiting cystatin;
l tumour; testis malformation; PCR primer; ss.
                                               Similarity
                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                 Toehoenen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98SE-00001617.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-SE000764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to amplify mouse
                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                     protein used
                                                                                                                                                                                                                                                                                                                        English.
                                 0,
                                               Score 15;
Pred. No.
                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          testatin
                                                                                                                                                                                                                                                                                                                                                                      for testis tumor diagnostics
                                                1.3e+03;
                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA
                                                               Length 15;
                                    Indels
                                    0
                                    Сарв
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THE PRICE OF SECTION O
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AAZ64910
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                      27-APR-1998;
18-SEP-1998;
25-FEB-1999;
23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                            Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavage; cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with interferon to treat HCV infection, other infectious diseases, autoimmune diseases, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence in the 3' non-core region. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 0
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                                                                                                                                                                                                                                                                      W09955847-A2
                                                                                                                                                                                                                                                                                                                         Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Substrate for HH ribozyme HCV.3-118 which cleaves HCV at nt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ64910 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 49; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel ribozymes for the treatment hepatitis C infection.
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25-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-APR-1998;
                                                                                                                                                            26-APR-1999;
                                                                                                                                                                                                                  04-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ64910;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 0; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              บบบบบบบบบบบบบบบบ 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                 98US-0083217P.
98US-0100842P.
99US-00257608.
99US-00274553.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0083217P.
98US-0100842P.
99US-00257608.
99US-00274553.
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                                                                                                                                                         99WO-US009027.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; 0 C; 0 G; 0 T; 15 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15; DB 1;
Pred. No. 1.3e+03;
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               [H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pavco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9418
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AAA4C502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with interferon to treat HCV infection, other infectious diseases, autoimmune
                                                    New DNA encoding endo-beta-mannanase from coffee, used e.g. in pharmaceutical, cosmetic or food compositions to hydrolyze polymannans.
                                                                                                                         Marraccini P,
                                                                                                                                                                               11-NOV-1998;
                                                                                                                                                                                                           28-OCT-1999;
                                                                                                                                                                                                                                                                 WO200028046-A1
                                                                                                                                                                                                                                                                                                                      Hydrolysis;
PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                             04-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA46502 standard; cDNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents the preferred target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 102; 123pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel ribozymes for the treatment hepatitis C infection.
                                                                                               WPI; 2000-399535/34.
                                                                                                                                                     (NEST )
                                                                                                                                                                                                                                      18-MAY-2000.
                                                                                                                                                                                                                                                                                            Coffea arabica
                                                                                                                                                                                                                                                                                                                                                               PCR primer used to amplify DNA encoding an endo-beta-mannanase.
                                                                                                                                                                                                                                                                                                                                                                                                                        AAA46502;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                     SOC PROD NESTLE SA.
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0; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mcswiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and cancer
                                                                                                                                                                                                                                                                                                                      polysaccharide; mannan; coffee; endo-beta-mannanase;
88.
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                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                         Rogers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15; DB 1;
Pred. No. 1.3e+03;
15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       μ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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PCR primers AAA46501-02 were used to amplify DNA encoding an endo-beta-

Page 32;

41pp;

French.

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Query Match
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The present primer was used to reverse transcribe human RNA, from which a CDNA sequence encoding a protein with heparanase catalytic activity was amplified. The heparanase (hpa) polynucleotide is useful in gene therapy, particularly in treating tumour, inflammation or autoimmunity. Particularly, the polynucleotide is useful in modulating the bioavailability of heparin-binding growth factors, cellular responses to heparin-binding growth factors (e.g. bFGF) and cytokines (e.g. interleukin (IL)-8), cell interaction with plasma lipoproteins, cellular susceptibility to certain viral and some bacterial and protozoa infections, or disintegration of neurodegenerative plaques. The polynucleotide is also useful in wound healing (e.g. thermal, chemical or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; heparanase; gene therapy; tumour; inflammation; autoimmunit heparin-binding growth factor; cytokine; neurodegenerative plaque; wound healing; infection; burn; angiogenesis; restenosis; atherosclerosis; inflammation; neurodegenerative disease; gerstmann-Straussler Syndrome; Creutzfeldt-Jakob disease; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                New polynucleotides encoding a polypeptide having heparanase activity, useful in wound healing and in gene therapy, particularly in treating tumor, inflammation, autoimmunity, neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-FEB-2000; 2000WO-US003542
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INSI-)
(HADA-)
(FRIE/)
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 I, Vlodavsky I,

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                                                                                                                                                                                                                                                           Page 44; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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Pred. No. 1.3e+
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .3e+03
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The invention provides modified nucleomonomers of specified formula and their pharmaceutically acceptable salts. The nucleomonomers are used as componers in oligomers, which are used in pharmaceutical compositions to consider in the composition of nucleic acid molecules including DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-ci infected cells. They are used in oligomers for gene regulation, antisense technology, diagnostic applications to detect target sequences in compositions or compositions and such as those containing pathogenic bacteria, funging and viruses, oncogenes, growth hormones and enzymes, to target genes or conclecules, receptor molecules, oytokines, oncogenes, growth factors and conditions, cardiovascular disorders, immune reactions, cancer, viral infections and bacterial infections (see AAA07786 for details of other nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair conglesses, triplexes or other forms of stable association. Sequences
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligomers used to inhibit expression of nucleic acids and regulation, antisense technology and diagnostics.
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Pred. No. 1.3e+03;
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     The invention provides modified nucleomonomers of specified formula and CC their pharmaceutically acceptable salte. The nucleomonomers are used as CC monomers in oligomers, which are used in pharmaceutical compositions to CC inhibit expression of nucleic acid molecules including DNA and RNA in CC ells such as bacterial, fungal, yeast, mammalian, cancer and virally-CC infected cells. They are used in oligomers for gene regulation, antisense CC technology, diagnostic applications to detect target sequences in CC included RNAs that encode enzymes, bormones, serum proteins, adhesion CC molecules, receptor molecules, cytokines, oncogenes, growth factors and CC interleukins associated with pathological conditions, cardiovascular disorders, immune reactions, cancer, viral CC conditions and bacterial infections (see AAA07786 for details of other CC uses for which the oligomers are suitable for). Oligomers comprising the CC target nucleic acid sequences, are physiologically when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and CC fidelity for target DNA sequences, are physiologically stable, non-toxic and CC single-or double-stranded target nucleic acid binding activity to form CC AAA07786-803 represent oligonucleotides forming a third strand along with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation; psoriasis; duplex; ss.
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                                                                                                                                                                                                                                          their pharmaceutically acceptable salts. The nucleomonomers are used as monomers in oligomers, which are used in pharmaceutical compositions to inhibit expression of nucleic acid molecules including DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-infected cells. They are used in oligomers for gene regulation, antiser technology, diagnostic applications to detect target sequences in biological samples such as those containing pathogenic bacteria, fungi and viruses, oncogenes, growth hormones and enzymes, to target genes or encoded RNAs that encode enzymes, hormones, serum proteins, adhesion
                                                                                 conditions, cardiovascular disorders, immune reactions, cancer, viral infections and bacterial infections (see AAA07786 for details of other uses for which the oligomers are suitable for). Oligomers comprising the nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair fidelity for target DNA sequences. The oligomers demonstrate significant
                                 single- or double-stranded target nucleic acid binding activity to form duplexes, triplexes or other forms of stable association. Sequences AAAO7820-834 represent sequences forming triplex oligomers
                                                                                                                                                                                                       molecules, receptor molecules, cytokines, oncogenes, growth factors and interleakins associated with pathological conditions such as inflammatory
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                                                                                                   conditions, cardiovascular disorders, immune reactions, cancer, viral infections and bacterial infections (see AAA07786 for details of other uses for which the oligomers ere suitable for). Oligomers comprising the nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair fidelity for target DNA sequences. The oligomers demonstrate significant single- or double-stranded target nucleic acid binding activity to form duplexes, triplexes or other forms of stable association. Sequences AAA07788-803 represent oligonucleotides forming a third strand along with
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regulation, antisense technology and diagnostics.
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The invention provides modified nucleomonomers of specified formula and their pharmaceutically acceptable salts. The nucleomonomers are used as compositions to compositions to inhibit expression of nucleic acid molecules including DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-compositions to detect target sequences in childing DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-compositions to detect target sequences in cological samples such as those containing pathogenic bacteria, funging biological samples such as those containing pathogenic bacteria, funging and viruses, oncogenes, growth hormones, serum proteins, adhesion cellecules, receptor molecules, cytokines, oncogenes, growth factors and interleukins associated with pathological conditions, cardiovascular disorders, immune reactions, cancer, viral conditions, cardiovascular disorders, immune reactions, cancer, viral conditions and bacterial infections (see AAA07786 for details of other cuses for which the oligomers are suitable for). Oligomers comprising the nucleomonomers exhibit increased duplex DNA stability when hybridizing to target mucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair fidelity for target DNA sequences. The oligomers demonstrate significant cubecase, triplexes or other forms of stable association. Sequences the duplex parameters.
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C;
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G;
                         Score 15; DB 1; 1
Pred. No. 1.3e+03;
                                                                                                                                        13 T; 2
                                                                                                                                        U; 0 Other;
                                                                Length 15;
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4464 TTTTTTTTTTTTTT 4478

N V

Mismatches

Indels

0

Gaps

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UTTTTTTTTTTTTU 15

4464 TYTTTTTTTTTTT 4478

Matches Query Match Best Local

13;

Conservative

<u>ب</u>

Mismatches

Indels

0,

Gaps

0

Local

Similarity

0.2%;

Score 15; DB 1; Length 15; Pred. No. 1.3e+03;

Sequence 15

BP; 0 A; 0 C; 0 G; 13 T; 2 U; 0 Other;

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RESULT 2707
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ID AAA
CC their pharmace processor of the components are used as C monomers in oligomers, which are used in pharmaceutical compositions to cinhibit expression of nucleic acid molecules including DNA and RNA in CC cells such as bacterial, fungal, yeast, mammalian, cancer and virally-clinifected cells. They are used in oligomers for gene regulation, antisense crechnology, diagnostic applications to detect target sequences in the biological samples such as those containing pathogenic bacteria, funging and viruses, oncogenes, growth hormones and enzymes, to target genes or cencoded RNAs that encode enzymes, hormones, serum proteins, adhesion condecules, receptor molecules, cytokines, oncogenes, growth factors and interleukins associated with pathological conditions such as inflammatory conditions, cardiovascular disorders, immune reactions, cancer, viral confictions, cardiovascular disorders, cancer, viral c
                                          the duplex sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 20; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified nucleomonomers, used in physiologically stable, oligomers used to inhibit expression of nucleic acids an regulation, antisense technology and diagnostics.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            psoriasis; duplex; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid sequence of ODN-j.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention provides modified nucleomonomers of specified formula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTTTTUTTTUTTTT 15
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RESULT 2708
AAA07799
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                                                                                                                                                                                          The invention provides modified nucleomonomers of specified formula and CC their pharmaceutically acceptable salts. The nucleomonomers are used as CC monomers in oligomers, which are used in pharmaceutical compositions to CC inhibit expression of nucleic acid molecules including DNA and RNA in CC cells such as bacterial, fungal, yeast, mammalian, cancer and virally-cc infected cells. They are used in oligomers for gene regulation, antisense CC technology, diagnostic applications to detect target sequences in CC biological samples such as those containing pathogenic bacteria, funging and viruses, oncogenes, growth hormones and enzymes, to target genes or CC encoded RNAs that encode enzymes, hormones, serum proteins, adhesion CC interleukins associated with pathological conditions such as inflammatory CC conditions, cardiovascular disorders, immune reactions, cancer, viral infections and bacterial infections (see AAA07786 for details of other cuses for which the oligomers are suitable for). Oligomers comprising the CC nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair C fidelity for target DNA sequences. The oligomers demonstrate significant customers, triplexes or other forms of stable association. Sequences or the properson oligomucleotides forming a third strand along with
                                                                                         Query Match
Best Local
                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA07799 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified nucleomonomers, used in physiologically stable, non-toxic oligomers used to inhibit expression of nucleic acids and in gene regulation, antisense technology and diagnostics.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               psoriasis; duplex; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid sequence of ODN-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 20; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-AUG-1998;
                                                                                                                                            Sequence 15 BP; 0 A; 0 C; 0 G; 14 T; 1 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-AUG-1999;
                                                                                                                                                                                duplex sequences
                                    4464
                                                                        14;
 1
                                                                                           Similarity
                   TTTTTTTTTTTTT 4478
 TITTTTTTTTTTTT 15
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NEBRASKA.
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                                                                                       0.2%;
                                                                                         Score 15; I
Pred. No. 1.
                                                                        Mismatches
                                                                                     DB 1; I
1.3e+03;
                                                                                                          Length 15
                                                                        Indels
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RESULT 2709

RESULT 2710

0

AAA07825

AAA07825 standard; DNA; 15 BP

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AAA07802

XX AAA0

XX AAA0

XX AAA0

XX AAA0

XX Nucl

XX Nucl

XX Nucl

XX Nucl

XX Nucl

XX PN W02

XX PN W02

XX O2-1

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XX PR 22-;

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XX Micl

XX PN W02

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                                                                                                                                                                                                                                                                                                                                                                                              CC The invention provides modified nucleomonomers of specified formula and cc their pharmaceutically acceptable salts. The nucleomonomers are used as components in oligomers, which are used in pharmaceutical compositions to cinhibit expression of nucleic acid molecules including DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-cc infected cells. They are used in oligomers for gene regulation, antisense technology, diagnostic applications to detect target sequences in cological samples such as those containing pathogenic bacteria, funging biological samples such as those containing pathogenic bacteria, funging conditions, receptor molecules, cytokines, oncogenes, to target genes or considered RNAs that encode enzymes, hormones, serum proteins, adhesion concerned RNAs that encode enzymes, hormones, serum proteins, adhesion contextions, cardiovascular disorders, immune reactions, cancer, viral conditions, cardiovascular disorders, immune reactions, cancer, viral conditions and bacterial infections (see AAA07786 for details of other cuses for which the oligomers are suitable for). Oligomers comprising the concile mucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair conditions, triplexes or other forms of stable association. Sequences can be the process of the forms of stable association. Sequences or the form of stable association. Sequences or the form of stable forming a third strand along with the conditions of the condit
                                                                                                                                                Best Loc
Matches
                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified nucleomonomers, used in physiologically stable, non-toxic oligomers used to inhibit expression of nucleic acids and in gene regulation, antisense technology and diagnostics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA07802 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gold B;
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                                                                                                                                                                                                                                                                                                    Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                              the duplex sequences
                                                                                                                                                                                       Local
                                              4464 TTTTTTTTTTTTTT 4478 :|||||||||
                                                                                                                                                l Similarity
UTTTTTTTTTTTT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     duplex; ss.
                                                                                                                                                                                                                                                                                                        BP; 0 A; 0 C; 0 G; 13 T; 2 U; 0 Other;
                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence of ODN-0
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                                                                                                                                                                                       Score 15; DB 1; Length 15; Pred. No. 1.3e+03;
                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense technology; leukemia;
cellular proliferation;
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                                                                                                                                                    Gaps
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                                                                  RESULT 2711
AAA07831
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                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                               Matches
                     AAA07831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Modified nucleomonomers, used in physiologically stable, non-toxicoligomers used to inhibit expression of nucleic acids and in gene regulation, antisense technology and diagnostics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;
                                                 AAA07831 standard; DNA; 15 BP
                                                                                                                                                                                                                                                           Sequence 15 BP; 0 A; 0 C; 0 G; 14 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 30; 42pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid sequence of a strand of triplex oligomer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUN-2000
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                                                                                                                                                4464 TTTTTTTTTTTTTT 4478
                                                                                                                                                                                               14;
                                                                                                                                _
                                                                                                                                                                                                               Similarity
                                                                                                                              TTTTTTTTTTTT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         duplex; triplex; ss.
                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                       represent sequences forming triplex oligomers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                       0.2%;
                                                                                                                                                                                           Score 15; DB
Pred. No. 1.3e
1; Mismatches
                                                                                                                                                                                                          DB 1;
1.3e+03;
                                                                                                                                                                                             0
                                                                                                                                                                                                                           Length 15
                                                                                                                                                                                             Indels
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                                                                                                                                                                                           Gaps
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 AAAO7803
ID AAAC
XX
AC AAAC
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DT 23-:
XX
DE Nuc:
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                                                                                                                                                                                                                                                                                                       RESULT 2712
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Nucleic acid sequence of ODN-p.
                                 23-JUN-2000
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(first entry)

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The invention provides modified nucleomonomers of specified formula and their pharmaceutically acceptable salts. The nucleomonomers are used as monomers in oligomers, which are used in pharmaceutical compositions to inhibit expression of nucleic acid molecules including DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-infected cells. They are used in oligomers for gene regulation, antisense technology, diagnostic applications to detect target sequences in the concentration of the containing pathogenic bacteria, fungicand viruses, oncogenes, growth hormones, serum proteins, adhesion molecules, receptor molecules, hormones, serum proteins, adhesion molecules, receptor molecules, cytokines, oncogenes, growth factors and conterleukins associated with pathological conditions such as inflammatory conditions, cardiovascular disorders, immune reactions, cancer, viral infections and bacterial infections (see AAAO7786 for details of other nucleic acid sequences, are physiologically stable, non-toxic and the pathological conditions associated with pathological conditions cancer, viral suses for which the oligomers are suitable for). Oligomers comprising the nucleic acid sequences, are physiologically stable, non-toxic and the pathological conditions and bacterial infections are physiologically stable, non-toxic and the pathological conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair fidelity for target DNA sequences. The oligomers demonstrate significant single- or double-stranded target nucleic acid binding activity to form duplexes, triplexes or other forms of stable association. Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gold
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AAA07803;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA07820-834 represent sequences forming triplex oligomers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 30; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modified nucleomonomers, used in physiologically stable, non-toxic oligomers used to inhibit expression of nucleic acids and in gene regulation, antisense technology and diagnostics.
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                                                                            AAA07803 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid sequence of a strand of triplex oligomer 16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYNE-) UNIV NEBRASKA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                4464 TTTTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                           TTTTTTTTTTTTT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 0 A; 0 C; 0 G; 14 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                            DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15; DB 1;
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 15;
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The invention provides modified nucleomonomers of specified formula and their pharmaceutically acceptable salts. The nucleomonomers are used as CC monomers in oligomers, which are used in pharmaceutical compositions to CC inhibit expression of nucleix acid molecules including DNA and RNA in CC cells such as bacterial, fungal, yeast, mammalian, cancer and virally-CC infected cells. They are used in oligomers for gene regulation, antisense tethnology, diagnostic applications to detect target sequences in CC diological samples such as those containing pathogenic bacteria, funging and viruses, oncogenes, growth hormones and enzymes, to target genes or CC encoded RNAs that encode enzymes, hormones, serum proteins, adhesion CC interleukins associated with pathological conditions, cardiovascular disorders, immune reactions, cancer, viral CC conditions and bacterial infections (see AAA07786 for details of other CC uses for which the oligomers are suitable for). Oligomers comprising the CC nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and collected, triplexes or other forms of stable association. Sequences cancer, triplexes or other forms of stable association. Sequences or the forms of stable association. Sequences with the duplex of an appresent oligomucleotides forming a third strand along with the content of the content of the stable association.
                                                                                                                                                                                                                                                                                                        AAA07834
                                                                                                                                                                                                                                                                                                                                        RESULT 2713
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 20; 42pp; English.
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Nucleomonomer; cancer; gene regulation; antisense technology; leukemia;
                                                               Nucleic acid sequence of a strand of triplex oligomer 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 0
                                                                                                                                      23-JUN-2000
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                                                                                                                                                                                                                                                                        AAA07834 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      duplex sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 0; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                     DNA; 15
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bred. No. 1.3e+03;
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Nucleic 23-JUN-2000 AAA07796

acid

sequence of ODN-i. (first entry)

Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;

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AAA07796 standard; DNA; 15 BP

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their pharmaceutically acceptable salts. The nucleomonomers are used as commonmers in oligomers, which are used in pharmaceutical compositions to inhibit expression of nucleic acid molecules including DNA and RNA in cells such as bacterial, fungal, yeast, mammalian, cancer and virally-cinfected cells. They are used in oligomers for gene regulation, antisense technology, diagnostic applications to detect target sequences in conjugated samples such as those containing pathogenic bacteria, funging and viruses, oncogenes, growth hormones and enzymes, to target genes or cencoded RNAs that encode enzymes, hormones, serum proteins, adhesion conjugated RNAs that encode enzymes, hormones, serum proteins, adhesion conjugated conditions, cardiovascular disorders, immune reactions, cancer, viral conditions, cardiovascular disorders, immune reactions, cancer, viral conditions, cardiovascular disorders, immune reactions, cancer, viral consistent nucleic acid sequences are suitable for). Oligomers comprising the nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair fidelity for target DNA sequences. The oligomers demonstrate significant conjugates, triplexes or other forms of stable association. Sequences and conjugates, triplexes or other forms of stable association. Sequences
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                                                               Matches 13;
                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention provides modified nucleomonomers of specified formula and their pharmaceutically acceptable salts. The nucleomonomers are used as monomers in olicomers, which are used in pharmaceutical compositions to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified nucleomonomers, used in physiologically stable, non-toxic oligomers used to inhibit expression of nucleic acids and in gene regulation, antisense technology and diagnostics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 30; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   psoriasis;
                                                                                                                                                Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYNE-) UNIV NEBRASKA
4464 TTTTTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   duplex; triplex; 88
                                                                                                                                                  BP;
                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0097712P
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                                                                                                                                                  0 A; 0 C; 0 G; 13 T; 2 U; 0 Other;
                                                                                   0.2%;
                                                         2
                                                                                   Score 15; DB 1;
Pred. No. 1.3e+03;
                                                                   Mismatches
                                                                                                         Length 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC their pharmaceutically acceptable salts. The nucleonomers are used as CC monomers in oligomers, which are used in pharmaceutical compositions to CC inhibit expression of nucleic acid molecules including DNA and RNA in CC cells such as bacterial, fungal, yeast, mammalian, cancer and virally-CC infected cells. They are used in oligomers for gene regulation, antisense CC technology, diagnostic applications to detect target sequences in CC biological samples such as those containing pathogenic bacteria, fungi CC and viruses, oncogenes, growth hormones and enzymes, to target genes or CC molecules, receptor molecules, cytokines, oncogenes, growth factors and CC interleukins associated with pathological conditions such as inflammatory CC conditions, cardiovascular disorders, immune reactions, cancer, viral CC infections and bacterial infections (see AAAO7786 for details of other CC uses for which the oligomers are suitable for). Oligomers comprising the nucleonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and components are into cells while maintaining stringent base pair
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Best Local (
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                                                                                                                   Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;
                                                                                                                                                                                                                   Nucleic acid sequence of ODN-m.
                                                                                                                                                                                                                                                                                     23-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fidelity for target DNA sequences. The oligomers demonstrate significant single- or double-stranded target nucleic acid binding activity to form duplexes, triplexes or other forms of stable association. Sequences AAA07788-803 represent oligonucleotides forming a third strand along with the contract of the contract of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention provides modified nucleomonomers of specified formula their pharmaceutically acceptable salts. The nucleomonomers are used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-246530/21.
                             Synthetic
                                                                                         psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                            AAA07800 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 0 A; 0 C; 0 G; 11 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 20; 42pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UTTTUTTUTTTU 15
                                                                                         duplex;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15; DB 1; Length 15; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  strand along with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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psoriasis;

duplex; ss.

WO200011013-A1

Nucleomonomer; cancer; gene regulation; viral infection; inflammatory response;

antisense technology; leukemia; cellular proliferation;

0

Nucleic acid sequence of ODN-f.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 2716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC cells such as bacterial, fungal, yeast, mammalian, cancer and virally-
CC infected cells. They are used in oligomers for gene regulation, antisense
CC technology, diagnostic applications to detect target sequences in
CC biological samples such as those containing pathogenic bacteria, fungi
CC and viruses, oncogenes, growth hormones and enzymes, to target genes or
CC encoded RNAs that encode enzymes, hormones, serum proteins, adhesion
CC molecules, receptor molecules, cytokines, oncogenes, growth factors and
CC interleukins associated with pathological conditions such as inflammatory
CC conditions, cardiovascular disorders, immune reactions, cancer, viral
CC infections and bacterial infections (see AAA07786 for details of other
CC uses for which the oligomers are suitable for). Oligomers comprising the
CC uncleomonomers exhibit increased duplex DNA stability when hybridizing to
CC target nucleic acid sequences, are physiologically stable, non-toxic and
CC fidelity for target DNA sequences. The oligomers demonstrate significant
CC fidelity for target DNA sequences. The oligomers demonstrate significant
CC fidelity for percent oligomers or other forms of stable association. Sequences
CC AAA07788-803 represent oligonucleotides forming a third strand along with
                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                           AAA07793 standard; DNA; 15 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modified nucleomonomers, used in physiologically stable, oligomers used to inhibit expression of nucleic acids an regulation, antisense technology and diagnostics.
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                                                                                                                                                                                       23-JUN-2000
                                                                                                                                                                                                                         AAA07793;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 20; 42pp; English.
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Pred. No. 1.3e+03;
2; Mismatches 0
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Best Local Similarity
Thes 0; Conserve
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                                                                                                                                                  Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;
                                                                                                                                                                                                      Nucleic acid sequence of ODN-k.
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                  02-MAR-2000
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Pred. No. 1.3e+03;
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conditions, cardiovascular disorders, immune reactions, cancer, viral infections and bacterial infections (see AAA07786 for details of other uses for which the oligomers resultable for). Oligomers comprising the nucleomonomers exhibit increased duplex DNA stability when hybridizing to target nucleic acid sequences, are physiologically stable, non-toxic and able to penetrate into cells while maintaining stringent base pair fidelity for target DNA sequences. The oligomers demonstrate significant single- or double-stranded target nucleic acid binding activity to form duplexes, triplexes or other forms of stable association. Sequences AAA07788-803 represent oligonucleotides forming a third strand along with
                                                                                                                                                                                                                                                                                                                   cells such as bacterial, fungal, yeast, mammalian, cancer and virally-infected cells. They are used in oligomers for gene regulation, antisense technology, diagnostic applications to detect target sequences in biological samples such as those containing pathogenic bacteria, fungi and viruses, oncogenes, growth hormones and enzymes, to target genes or encoded RNAs that encode enzymes, hormones, serum proteins, adhesion molecules, receptor molecules, cytokines, oncogenes, growth factors and interleukins associated with pathological conditions such as inflammatory conditions associated with pathological conditions such as inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               their pharmaceutically acceptable salts. The nucleomonomers are used a monomers in oligomers, which are used in pharmaceutical compositions to inhibit expression of nucleic acid molecules including DNA and RNA in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified nucleomonomers, used in physiologically stable, non-toxic oligomers used to inhibit expression of nucleic acids and in gene regulation, antisense technology and diagnostics.
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                                                  the duplex sequences
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Query Match
Best Local
Similarity 0; Conserv
           0.2%;
          Score 15; DB 1; Length 15; Pred. No. 1.3e+03;
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BP; 0 A; 0 C; 0 G; 0 T; 15 U; 0 Other;

Sequence 15

δ Matches 4464 TITTTTTTTTTTTT 4478 Conservative 15; Mismatches Indels <u>,</u>

Gaps

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AAA07788 standard; DNA; 15
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Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;

Nucleic acid sequence of ODN-a.

23-JUN-2000

(first entry

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02-MAR-2000.
                        Synthetic
                               psoriasis; duplex; ss.
   20-AUG-1999;
                 WO200011013-A1
   99WO-US019029
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Cl infected cells. They are used in oligomers for gene regulation, antisense
CC technology, diagnostic applications to detect target sequences in
CC biological samples such as those containing pathogenic bacteria, fungi
CC and viruses, oncogenes, growth hormones and enzymes, to target genes or
CC encoded RNAs that encode enzymes, hormones, serum proteins, adhesion
CC molecules, receptor molecules, cytokines, oncogenes, growth factors and
CC interleukins associated with pathological conditions such as inflammatory
CC conditions, cardiovascular disorders, immune reactions, cancer, viral
CC uses for which the oligomers are suitable for). Oligomers comprising the
CC uses for which the oligomers are suitable for) conditions non-toxic and
CC able to penetrate into cells while maintaining stringent base pair
CC fidelity for target DNA sequences. The oligomers demonstrate significant
CC single- or double-stranded target nucleic acid binding activity to form
CC duplexes, triplexes or other forms of stable association. Sequences
CC AAA07788-803 represent oligonucleotides forming a third strand along with
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                                                                                                                                                                                                                 Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation;
                  22-AUG-1998;
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                                                                                                                                                                Synthetic.
                                                                                                                                                                                              psoriasis; duplex; ss.
                                                                                                                                                                                                                                                                      Nucleic acid sequence of ODN-d
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                                                                                                                                                                                                                                                                                                                                                                                  AAA07791 standard; DNA; 15
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Pred. No. 1.3e+
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cc cells such as bacterial, fungal, yeast, mammalian, cancer and virally-
cl infected cells. They are used in oligomers for gene regulation, antisense
ct technology, diagnostic applications to detect target sequences in
cc biological samples such as those containing pathogenic bacteria, fungi
cc and viruses, oncogenes, growth hormones and enzymes, to target genes or
cc encoded RNAs that encode enzymes, hormones, serum proteins, adhesion
cc molecules, receptor molecules, cytokines, oncogenes, growth factors and
cc interleukins associated with pathological conditions such as inflammatory
cc conditions, cardiovascular disorders, immune reactions, cancer, viral
cc infections and bacterial infections (see AAA07786 for details of other
cuses for which the oligomers are suitable for). Oligomers comprising to
cuses for which the oligomers are suitable for) conditions, cancer,
cc nucleomonomers exhibit increased duplex DNA stability when hybridizing to
carget nucleic acid sequences, are physiologically stable, non-toxic and
cc able to penetrate into cells while maintaining stringent base pair
cf iddelity for target DNA sequences. The oligomers demonstrate significant
cc single- or double-stranded target nucleic acid binding activity to form
duplexes, triplexes or other forms of stable association. Sequences

AAA07788-803 represent oligonucleotides forming a third strand along with
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                                                     duplex sequences
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Query Match Sequence 15 BP; 0 A; 0 C; 0 G; 11 T; 4 U; 0 Other, 0.2%; Score 15; DB 1; Pred. No. 1.3e+03; Length 15;

Ś 밁 4464 TTTTTTTTTTTTTT 4478 1

Best Local Similarity Matches 11; Conserv

Conservative

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Mismatches

<u>.</u>

Indels

<u>,</u>

Gaps

0

AAA07801 standard; DNA; 15 BP

Nucleic acid

23-JUN-2000 AAA07801;

(first entry)

sequence of ODN-n.

Nucleomonomer; cancer; gene regulation; antisense technology; leukemia; viral infection; inflammatory response; cellular proliferation; psoriasis; duplex; ss.

Synthetic

WO200011013-A1

02-MAR-2000

20-AUG-1999; 99WO-US019029.

DIX AX FX BX X OX X BX FX FX FX BX BX S X BX S X BX S 22-AUG-1998; 98US-0097712P

(UYNE-) UNIV NEBRASKA

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RESULT 2721
AAA62350
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oligomers used to inhibit expression of nucleic acids and in
regulation, antisense technology and diagnostics.
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                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                   Oligonucleotide
04-JUL-2000
                                                                                                                                                                                                                                                                                Conformationally-locked oligonucleotide; antisense inhibitor; bicyclic sugar nucleoside analogue; gene probe; ds.
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                                                                                                                                                          /*tag=
/mod_ba
                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                        'note=
                                                                     note=
                                                                                                                                                                                                                                                                                                                               containing 3'-C-amino-5'(S)-C,3'-N-ethanothymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     73.3%;
                                                          base= OTHER
:= "3'-C-amino-5'(S)-C,3'-N-ethanothymidine'
                                                                                                                                                          base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
                                                                                                                                        "3'-C-amino-5'(S)-C,3'-N-ethanothymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15; DB 1;
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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modified_base

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/*tag= c /mod_base=

note=

3,

-C-amino-5' OTHER

(R) -C, 3'-N-ethanothymidine"

modified_base

/mod_base= OTHER /note= "3'-C-amino-5'(R)-C,3'-N-ethanothymidine"

/*tag= __ /mod_base= (

OTHER

(R)-C,3'-N-ethanothymidine"

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RESULT 2722
AAA62347
PRINTER TO THE PRINTE
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Matches 15; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is an oligonucleotide containing 3'-C-amino-5'(S)-C,3'-N-ethanothymidine, a bicyclic-sugar nucleoside. All nucleotides in the sequence were incorporated by phosphoramidite chemistry using a DNA synthesiser. Bicyclic sugar nucleosides are conformationally restricted 3',5'-bridged nucleosides which can be used as building blocks for oligonucleotides. Oligonucleotides can be produced that have certain, desired, geometrical shapes and entropy advantages. They may have superior hybridisation to DNA and RNA, and excellent biological stability. The conformationally-modified oligonucleotides may be useful as antisense inhibitors of gene expression or as gene probes, and may therefore be used in antisense therapeutics or gene-specific diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New conformationally restricted 3',5'-bridged nucleosides and oligonucleotides useful as antisense therapeutics or as gene-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 20; Col 16; 10pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-451496/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conformationally-locked oligonucleotide; bicyclic sugar nucleoside analogue; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide #3 containing 3'-C-amino-5'(R)-C,3'-N-ethanothymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA62347 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4464 TITTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-00309742
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense inhibitor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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AAA62348
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                     The present sequence is an oligonucleotide containing 3'-C-amino-5'(R)-C,3'-N-ethanothymidine, a bicyclic-sugar nucleoside. All nucleotides in the sequence were incorporated by phosphoramidite chemistry using a DNA synthesiser. Bicyclic sugar nucleosides are conformationally restricted 3',5'-bridged nucleosides which can be used as building blocks for oligonucleotides. Oligonucleotides can be produced that have certain, desired, geometrical shapes and entropy advantages. They may have superior hybridisation to DNA and RNA, and excellent biological stability. The conformationally-modified oligonucleotides may be useful as antisense inhibitors of gene expression or as gene probes, and may therefore be used in antisense therapeutics or gene-specific diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
         modified_base
                                                                     Conformationally-locked oligonucleotide; antisense inhibitor; bicyclic sugar nucleoside analógue; gene probe; ds.
                                                                                                                                     06-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New conformationally restricted 3',5'-bridged nucleosides and oligonucleotides useful as antisense therapeutics or as gene-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-JUL-2000
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                                                                                                           Oligonucleotide #4 containing 3'-C-amino-5'(R)-C,3'-N-ethanothymidine
                                                                                                                                                              AAA62348;
                                                                                                                                                                                     AAA62348 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-451496/39.
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                                                                                                                                                                                                                                                                                                  15; Conser
                                                                                                                                                                                                                                                                      TTTTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                                   15 BP; 0 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC
                                                                                                                                   (first
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             Location/Qualifiers
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/mod_ba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note = "3'-C-amino-5'(R)-C,3'-N-ethanothymidine
                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                  0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                _base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                base= OTHER
                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "3'-C-amino-5'(R)-C,3'-N-ethanothymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ase= OTHER
"3'-C-amino-5'(R)-C,3'-N-ethanothymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                  k; Score 15; DB
k; Pred. No. 1.3
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                  15 T; 0
                                                                                                                                                                                                                                                                                                                                                   U; 0 Other;
                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                             1.3e+03;
                                                                                                                                                                                                                                                                                                                         Length 15;
                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                  0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                      the sequence were incorporated by phosphoramidite chemistry using a DNA synthesiser. Bicyclic sugar nucleosides are conformationally restricted 3',5'-bridged nucleosides which can be used as building blocks for oligonucleotides. Oligonucleotides can be produced that have certain, desired, geometrical shapes and entropy advantages. They may have superior hybridisation to DNA and RNA, and excellent biological stability. The conformationally-modified oligonucleotides may be useful as antisense inhibitors of gene expression or as gene probes, and may therefore be used in antisense therapeutics or gene-specific diagnostics
                      modified_base
                                                                                 modified_base
                                                                                                                        Synthetic
                                                                                                                                                                       Oligo dT15
                                                                                                                                                                                                  31-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is an oligonucleotide containing 3'-C-amino-5'(R)-C,3'-N-ethanothymidine, a bicyclic-sugar nucleoside. All nucleotides in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-JUL-2000
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                                                                                                                                                 Hybridisation probe; DNA cleavage; double-helix; oncogene; ss.
                                                                                                                                                                                                                           AAH20308;
                                                                                                                                                                                                                                                 AAH20308 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 20; Col 15; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotides useful as antisense therapeutics or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New conformationally restricted 3',5'-bridged nucleosides and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-MAY-1999;
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15; Conserva
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                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                  at C-5"
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/mod_base= OTHER
/note= "3'-C-amino-5'(R)-C,3'-3'-N-ethanothymidine"
                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note=
         /*tag= b
                                                 /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'mod_base= OTHER
'note= "3'-C-amino-5'(R)-C,3'-3'-N-ethanothymidine"
                                                            mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                               "Optionally thymidine has EDTA covalently attached
                                                                                                                                                                                                                                                   15
                                                                                                                                                                       probe.
                                                                                                                                                                                                                                                                                                                15
                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                 ,;
                                                                                                                                                                                                                                                                                                                                                                              Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 15;
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                                                                                                                                                                                                                                                                                                                                                                 Indels
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-OCT-1987;
16-NOV-1990;
12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to hybridisation probes which target a specific sequence within a large double-helical nucleic acid. The probe is complementary to the target sequence and contains at least one nucleotide with an attached molecule that is able to cleave double-helical DNA e.g. EDTA-Re(II) (ethylenediaminetetrascetic acid-iron complex). The probes where the attached molecule is a label or compound that alters gene expression, are used for specific detection and/or cleavege of double-helical DNA, e.g. for diagnosis, for treatment of disease (particularly caused by viruses, genetic defects or oncogenes), for chromosomal analysis, and for the isolation and mapping of genes. The present sequence represents probe of the invention used in an example illustrating how the probe binds to and cleaves double stranded DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New hybridization probe for specific triplex formation with large double helices, useful e.g. for site-specific diagnostic cleavage, contains attached functional residue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-AUG-1998;
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                                                                   Synthetic
                                                                                             ODN-MGB-LF; oligonucleotide; minor groove binder; latent fluorophore; hybridisation; detection; fluorescence; probe; ss.
                                                                                                                                        Oligonucleotide portion of ODN-MGB-LF conjugate.
                                                                                                                                                                                                                            AAF30882 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-342909/36.
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              03-MAY-2001
                                         WO200131063-A1
                                                                                                                                                                    09-JUL-2001
                                                                                                                                                                                                AAF30882;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Fig 3B; 20pp; English.
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                                                                                                                                                                                                                                                                                                                          4464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PB,
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                                                                                                                                                                                                                                                                                                                  TTTTTTTTTTTTT 4478
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                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                  (first entry)
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90US-00614205.
93US-00152250.
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/note= "Optionally thymidine has EDTA covalently attached
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note= "Optionally thymidine has EDTA covalently attached
                                                                                                                                                                                                                                                                                                                                                                                                             A; 0 C; 0 G; 15 T; 0
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                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
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                                                                                                                                                                                                                                                                                                 15
                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                      Score 15; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                             U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                    1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      one nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conjugate of oligonucleotide, useful for detecting specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 58; 105pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-328656/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dempcy RO,
                                          DE10051726-A1
                                                                                                                                                  modified_base
                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                            clinical diagnostic;
                                                                                                                                                                                                                                                                                                    Cross-linked vinyl acetate copolymer carrier material; AIDS treatment;
                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide b) for solid phase synthesis of oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                            31-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH20511;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH20511 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mismatch discrimination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-OCT-2000; 2000WO-US029786
                                                                                                                                                                                                                                                                                  phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fluorescence
of 8.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPOC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4464 TTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Afonina IA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9905-00428236.
                                                   /*tag= a
/mod_base= OTHER
/mote= "Phosphorothioate deoxynucleotides"
                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                solid phase synthesis; modified oligonucleotide;
ic; cancer treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTT 4478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15; DB 1; Length 15; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      minor groove binder and latent fluorophore, nucleic acids, e.g. for single-nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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10-MAY-2001.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 15
                                                                                                                                                                    Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; lethyosis; serborrhoea; ruba; keratosis; neoplasia; sleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                      30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                AAF49041 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Intermediate for oligonucleotide synthesis comprises partially hydrolysed cross-linked vinyl acetate copolymer loaded with nucleotide derivative.
                            21-JUN-2000; 2000WO-AU000693.
                                                                                           WO200078341-A1
                                                                                                                           Homo sapiens
                                                                                                                                                         hyperneovascular condineovascular condi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seliger H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-OCT-2000;
                                                                                                                                                                                                                                                                                                    IGF-I oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention describes a novel chemical product comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001-336414/36
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                                                                                                                                                                                                                                                                                                                                  (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 A;
                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                    entry)
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Pred. No.
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. 1.3e+03;
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21-JUN-1999;

99US-0140345P

Wraight

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Werther GA,

Edmondson

(MURD-) 21-JUN-1999; 21-JUN-2000; 28-DEC-2000. WO200078341-A1.

MURDOCH CHILDRENS RES

INST

2000WO-AU000693

99US-0140345P

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AAF45344/c
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                               Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; kelo skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasi IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris growth factor mediated cell proliferation; ichthyosis; serborrhoea; rukeratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic diseas hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for ameliorating the effects skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Pactor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MURD-) MURDOCH CHILDRENS RES
                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                                                           AAF45344 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                      neovascular condition
                                                                                                                                                                                                                                                                                                                                    IGFBP2 oligonucleotide #183.
                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                AAF45344;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wraight CJ,
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                                                                                                                                                                         sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
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hes 0;
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                                                                                                                                                                                                                                                             pityriasis;
s; pilaris;
                                                                                                                                                                                                                                   disease;
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                                                                                                                                                                                                                                                                                           keloid,
                                                                                                                                                                                                                                                  ruba
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AAH18783/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           c skin disorders. The method comprises contacting the skin with an cantisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an coligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood coesses is a condition of the retina, cancers of the growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood conditions are conditionally the condition of the skin, and the condition of the skin cancers of the growth factor-mediated malignancies, other sclerotic conditions of the skin cancers of the growth factor-mediated malignancies, other sclerotic conditions of the skin cancers of the growth factor-mediated malignancies.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8783/c
AAH18783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-041421/05
                                                                                                                             Chew
                                                                                                                                                                                                                                                                                                                                                   paternity test;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH18783;
                                            Polynucleotide comprising novel single interleukin-4 gene for use in studying
                                                                                                                                                                                                                                                        05-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation.
                             interleukin-4,
                                                                                                                                                                                                                       28-SEP-2000; 2000WO-US026608
                                                                                                                                                                                                                                                                                      WO200123404-A1
                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                  inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7414
                                                                                             2001-316132/33.
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                                                                                                                                                                                                                                                                                                                                                                                                                               IL4 allele-specific primer SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                 interleukin-4; IL4; single nucleotide polymorphism; SNP; atopy;
matory disorder; immune disorder; population diversity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15;
                                                                                                                                                            GENAISSANCE PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6; Page 35; 201pp; English
                                                                                                                             Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGCAGCAGCAGCAGC 7428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention relates to a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                             Y,
                               in developing drugs,
                                                                                                                                                                                           99US-0156825P.
                                                                                                                                                                                                                                                                                                                                                                  forensic test; cytokine; chromosome 5q31.1; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                             Denton
                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
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Pred. No.
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                               diagnosis
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                                              nucleotide perpression,
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1.3e+03;
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                               and
                                                                                                                           Stephens
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                                              polymorphisms, function of
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                               treatment
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein, cDNA and gene of human interleukin-4 (IL4). The coding sequences for this protein contain single nucleotide polymorphisms (SNPs) which may be associated with differences in susceptibility to atopy, inflammatory and immune diseases and different drug responses. They may also be used in applications such as forensic and paternity testing and studying population diversity and anthropological lineage. The IL4 gene is found on human chromosome 5q31.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Page 16; 71pp; English.
                                                                                                        WPI;
                                                                                                                           Uhlmann
                                                                                                                                                              14-MAR-1994;
08-MAR-1995;
                                                                                                                                                                                            08-MAR-1995;
                                                                                                                                                                                                               04-JUL-2001.
                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                    Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                     peptide nucleic acid; ss.
                                                                                                                                                                                                                                                                                                                                                                              antiviral; hepatotropic; vasotropic; antisense innibition; r
integrin; cell-cell adhesion; cancer; restenosis; stability;
                                                                                                                                                                                                                                                                                                                                                                                                 Polyamide-oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                   PNA-forming
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                                                                                                                                             (AVET ) AVENTIS PHARMA DEUT GMBH
                                                                                                                                                                                                                                  EP1113021-A2
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                                                                                                        2001-591267/67.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                          Breipohl
                                                                                                                                                                                            2001EP-00104012
                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                               94DE-04408528.
95EP-00103332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 A;
                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                    /*tag= b
/mod_base= OTHER
/note= "t-hex"
                                                                                                                                                                                                                                                                                         note=
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                                                                                                                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         "t-but"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5813
                                                                                                                                                                                                                                                                                                                                                                                       e derivative; anticancer; vasotropic; antisense in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             먥
                                                                                                                                                                                                                                                                                                    OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                    #7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                        inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                  antiproliferative;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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This invention describes novel polyamide-oligonucleotide derivatives (I) and their physiologically acceptable salts of formula $F(\{DNA\}-Li\}$ g(PNA-Li) g(PNA-Li) g(PNA) g(PNA)

Example

Page 40;

54pp; German.

New DNA-peptide nucleic acid chimeras, useful e.g. as antisense agents $\dot{}$ for treating e.g. cancer, also as diagnostic probes and primers.

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the invention have anticancer, antiproliferative, antiviral, hepatotropic compression by antisense, ribozyme, sense, or triple-helix methods, or by binding to proteins (aptamers). (I) are used for the inhibition of gene companies of the proteins (aptamers). (I) are used for treating diseases caused by viruses (human immune deficiency, herpes simplex, influenza, vesicular storativis, hepatitis B or papilloma), or mediated by integrins or cell-cell adhesion reactions, for treating cancer, or for inhibiting crestenosis, particularly as antisense reagents. They are also useful in the heterogeneous or homogeneous assays, as primers or probes, particularly where the target is amplified before being detected by hybridization, for the increased affinity for complementary strands and better stability in the increased affinity for complementary strands and better stability in the increased affinity for complementary strands and better stability in the increased affinity conventional peptide nucleic acids (PNA), but lack the disadvantages, i.e. have improved cellular uptake, do not aggregate in aqueous solution, and have reduced affinity for purification complements, reduced cytotoxicity, better sequence specificity. They are more active than either DNA or PNA oligomers. When used as probes, (I) show different responses to base-pair mismatches in the DNA and PNA segments, allowing better discrimination between pathogenic and nongenetary better sequence specificaty. They are concepted to eliminate RNA or DNA primers. The DNA component allows to be used to eliminate RNA or DNA primers. The DNA component allows of additional reactions not possible with NA H49208-AAH49208 control of the proporated into a gene. AAH49208-AAH49208 control of the proporated into a gene. AAH49208 control of the proporated to the proporated to the proporated to the proporated to t
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Matches
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                                                                                                                                                            20-SEP-2000; 2000US-00666390.
                                                                                                                                                                                                                                                                                                                         21-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                   US2002034810-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gallus gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Heparanase; catalytic; cytostatic; antiviral; antibacterial; enzyme; anti-protozoan; neuroprotective; heparin; hpa; chicken; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chicken heparanase (hpa) cDNA cloning oligo dT(15) primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         may be incorporated into a gene. AAH49208-AAH49264 represent oligonucleotides used to illustrate the method of the invention
                                                                                                                                                                                                                                           16-AUG-2001; 2001US-00930218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JUL-2002
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                                                                          INSIGHT STRATEGY & MARKETING LTD
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Vlodavsky I, Michal I,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 T; 0 U; 0 Other;
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Nucleic acid encoding avian and reptile heparanase to treat various heparin-related disorders and the

polypeptide is useful signal peptide is

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  secreted proteins in heterologous expression systems. The encoded polypeptides can be used to prevent tumour anglogenesis, metastasis and invasion, and to intervene with pathologies associated with impaired heparin-binding growth factors, cellular responses to heparin-binding growth factors and cytokines, cell interaction with plasma lipoproteins, cellular susceptibility to viral, protozoa and bacterial infections or disintegration of neurodegenerative plaques. The present sequence represents a chicken heparanase (hpa) cDNA cloning oligo dT(15) primer
          separation of nucleic acid molecules and as probes, primers, linkers adapters and antisense agents on solid supports. Modifications enhan their use as capture and detection probes e.g. by the incorporation biotin, digoxigenen, radioisotopes, fluorescent labels such as fluorescein and reporter molecules such as alkaline phosphatase. The also used for enhancing or inhibiting the activity of an enzyme or cellular activity. The compounds are stable to nucleases and proteas
                                                                                                                                           This invention relates to oligonucleotide analogues comprising
                                                                                                                                                                                                          Oligonucleotides analogs useful in detection, of nucleic acid molecules, comprise monomers,
                                                                                                                                                                                                                                                                                                                                                                          14-MAR-2000;
30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein nucleic acid molecule; PNA; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated avian and reptile nucleic acid, encoding a polypeptide with heparanase catalytic activity. The signal peptide of the nucleic acid can be used to express membrane-associated or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      proceins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful in production of membrane-targeted or secreted recombinant
                                                                                                                                                                          Example 20; Page 123; 197pp; English.
                                                                                                                                                                                                                                                           WPI; 2002-041177/05.
                                                                                                                                                                                                                                                                                          Chakhmakhcheau O,
                                                                                                                                                                                                                                                                                                             Efimov V,
                                                                                                                                                                                                                                                                                                                                                                                                                        13-MAR-2001; 2001WO-US008111.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-SEP-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of oligomer # 10 used to compare mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUN-2002
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                                                                                                                             acid molecule (PNA)
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                                                                                                                                                                                                                                                                                                         Fernandez J,
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2000US-0250334P.
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                                                                                                                                                                                                                                                                                          Buryakova A,
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100.0%; Pr/
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry
                                                                                                                                                                                                                                                                                        Archdeacon D, A:
kova A, Choob M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    isolated avian and reptile nucleic acid, heparanase catalytic activity. The signal
                                                                                                                          monomer. They are used in the
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Pred. No.
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1.3e+03;
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M, Hondorp
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                                                                                                                                                                                                          and purification oligomers.
                                                                                                                            ising a protein detection and
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                                                                                                           linkers,
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RESULT 2733
AAL49453/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   backbone of PNAs is resistant to both nucleases and proteases. PNAs bind nucleic acid molecules with greater affinity than DNA or RNA concentration. The compounds are relatively simple to synthesize and are used in a wide variety of applications. This sequence represents a DNA oligomer which is used to represent the effect of single base mismatches
The present invention relates to a method of detecting mutations in a nucleic acid by amplifying the nucleic acid to produce a double-stranded amplicon, in vitro transcription and translation of this amplification, and detection of the translated protein. The primers used for amplification are designed to produce an amplicon that is translatable and allows differentiation between translation products of wild-type and mutated nucleic acids. The method is used to detect mutations in tumour suppressor genes, for (early) diagnosis, monitoring and characterisation of tumours (especially of bladder and intestines) and in the germ line (using nucleic acids from embryos or blood cells). A new multi-tag vector is used to detect or verify the reading frame of a nucleic acid cloned in it, and to determine the suitability of detectable peptides for analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mutation detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL49453
                                                                                                                                                                                                             Detecting mutations in nucleic acid, useful for diagnosis and characterization of tumors, by amplification, in vitro transc
                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-2002.
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                                                                                                                                                                       Claim 11; Fig
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                                                                                                                                                                       5; 62pp;
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cancer;
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                                                                                                                                                                                                 detection
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Pred. No.
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                                                                                                                                                                                                              in vitro transcription
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RASUUT 2734

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                     nucleic acid by amplifying the nucleic acid to produce a double-stranded amplicon, in vitro transcription and translation of this amplicon, and detection of the translated protein. The primers used for amplification are designed to produce an amplicon that is translatable and allows differentiation between translation products of wild-type and mutated nucleic acids. The method is used to detect mutations in tumour suppressor genes, for (early) diagnosis, monitoring and characterisation of tumours (especially of bladder and intestines) and in the germ line (using nucleic acids from embryos or blood cells). A new multi-tag vector is used to detect or verify the reading frame of a nucleic acid cloned in it, and to determine the suitability of detectable peptides for analysis and/or purification of a recombinant protein, expressed from a sequence cloned in the vector. The present sequence encodes a tag peptide and was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting mutations in nucleic acid, useful for characterization of tumors, by amplification, in translation, then protein detection.
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                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Fig 5; 62pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kahmann
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                                                                                                                                                                                                                                                                                                                                                                  present invention
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DB; AAO19056.
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cancer;
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Pred. No.
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1.3e+03;
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n vitro transcription and
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                                                                                                                                                                                                                      The present invention relates to new human pre-adipose cell lines capable to differentiate to white adipose cells, exhibiting essentially the same cellular properties of normal white adipose cells. The human pre-adipose cell lines are useful for the identification of substances controlling the regulation of lipid uptake and release by human white adipocytes, and substances controlling the differentiation of preadipocytes into mature adipocytes. They are useful for screening compounds capable to regulate the secretion of any metabolites or hormones from human white adipocytes. Sequences of the invention are useful for developing drugs, food ingredients and supplements against obesity, diabetes and cardiovascular diseases. The present DNA sequence is a reverse transcription (RT)-PCR primer which is used for the expression of adipocytes in differentiated immortalised human preadipose cells. This primer is used in the cardiocal sequence is a reverse transcription of the cardi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human pre-adipose cell line capable of differentiating to adipose cells, useful in developing drug, food ingredients, and supplements
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pre-adipose cell line; white adipocyte; food ingredient; diabetes; cardiovascular disease; reverse transcription;
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                                                                                                                                                                                                         the exemplification of the invention
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                          TTTTTTTTTTTTT 4478
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                                                                                                  Score 15;
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                                                                                Mismatches
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RESULT 2736
AAD22531/c
                        used to decrease the expression of oncogenes and thereby decrease the expression of cancer cells which rely upon oncogene expression for their phenotypic and pathological properties. The oligoribonucleotides are also used for increasing the effectiveness of antisense oligonucleotide targetted to a gene associated with a disease or a condition in an animal. To alter gene transcription and/or translation for any gene or gene segment responsible for expression, to inhibit viral reverse transcriptase, to inhibit the expression of leukaemia virus, hepatitis virus, oncogenes and human immunodeficiency virus. The present sequence is retroviral reverse transcriptase inhibitor DNP-poly [A] RNA fragment which is used in the transcriptase inhibitor DNP-poly [A] RNA fragment which is used in the treatment of moloney murine leukaemia virus (MULV) in mammals. (Updated on 07-AUG-2003 to correct OS field.) (Updated on 29-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-2003
07-AUG-2003
12-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to derivatised antisense oligoribonucleotides with enhanced membrane permeability and stability. The derivatised antisense oligoribonucleotide complementary to a sequence of nucleotides found in virus or a cell is useful for inhibiting e.g., viral reverse transcriptase. Derivatised antisense oligoribonucleotide is conjugated at the 2'-O position with a hydrophobic carrier reagent containing a poly substituted phenyl compound. The derivatised oligoribonucleotides are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Derivatized antisense oligoribonucleotide useful to inhibit e.g. viral reverse transcriptase comprises at the 2'-O position of the oligoribonucleotide, a hydrophobic carrier reagent containing a poly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               substituted phenyl compound.
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23-FEB-1994;
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hepatitis virus; human immunodeficiency virus; retroviral; DNP-poly [A];
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94US-00200650.
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OS field)
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                                                                   δ
                                                                                                                                                                                                                                                                                                                                                        The present invention describes a vector (I) comprising operably linked CC recipient cell, preferably in bacteria such as Escherichia coli; (b) clearly send cetable marker region capable of being expressed in the recipient cell; and (c) a chimeric DNA construct comprising: (i) promoter or promoter cell region capable of being expressed in the recipient cell cell or by prokaryotic RNA polymerase; (ii) first, second, third and cell courth recombination sites, (iii) 3' transcription terminating and cell fourth recombination sites, or the second and third recombination sites or the first and cell fourth recombination sites, do not recombination site, or the third and cell fourth recombination sites, do not recombination sites, or the third and cell fourth recombination sites, do not recombination sites, or the third and cell fourth recombination sites, do not recombination sites, or the third and cell sequences. The vectors can also be used to convert a DNA fragment cell into an inverted repeat structure. Plants transformed with a vector from the present invention can be used in a conventional breeding scheme to conventional breeding scheme to conventional material sequence a celds. The present sequence characteristics or to introduce a celds. The present sequence represents an acceptor vector mucleotide cells. The present sequence represents an acceptor vector mucleotide cells.
                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New vectors comprising operably linked DNA fragments having an origin of replication, a selectable marker and a chimeric DNA construct, useful for silencing target nucleic acids and for producing large amounts of double-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-JAN-2001; 2001US-0264067P.
29-NOV-2001; 2001US-0333743P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 14; Page 74; 104pp; English.
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                                                                                                                                                                                                                                                                         Sequence 15 BP;
                                                                                                                                                                          Local Similarity
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                                                                   TITITITITITIT 4478
                                                                                                                                      Conservative
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                                                                                                                                                                      100.0%;
                                                                                                                                                                                                          0.2%;
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                                                                                                                                      0
                                                                                                                                                                          Score 15;
Pred. No.
                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                             U; 0 Other;
                                                                                                                                                                                                             DB 1;
                                                                                                                                                                             1.3e+03
                                                                                                                                          0
                                                                                                                                                                                                          Length 15;
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                                                                                                                                          Indels
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ABXO0240

ABXO0240

AC ABXO

XX ABXO

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XX Heps

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                                                                                                                   Matches
                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therapy; type I interferon; interferon alpha; interferon beta; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                             of the substrate sequences defined in the specification. The HCV ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV ribozymes are also useful for treating a condition associated with HCV infection in conjunction with one or more other drug therapies, particularly type I interferon, especially interferon alpha, beta or gamma or consensus interferon. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to enzymatic nucleic acids which specifically cleave RNA derived from Hepatitis C virus (RCV). The enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin (HH) motif where the binding arms comprise sequences complementary to of the substrate sequences defined in the specification. The HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BLAT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABX00240 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABX00240;
                                                                                                                                                                                                                                   Sequence 15 BP; 0 A; 0 C; 0 G; 0 T; 15 U; 0 Other;
                                                                                                                                                                                                                                                                                              seqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                                                                                                                                                                          sequence represents a substrate for a HCV hammerhead (HH) ribozyme. No Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 21; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-617759/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2002082225-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PAVC/)
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                                                       4464 TTTTTTTTTTTTTT 4478
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PAVCO P A.
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MCSWIGGEN J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MACEJACK D
                                                                                                               Similarity 0; Conserv
Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gamma; consensus interferon; hepatotropic; antiinflammatory; hammerhead ribozyme; HH ribozyme; ss.
                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-00274553.
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                                                                                                                                               0.0%;
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                                                                                                                                               Pred.
                                                                                                                                                  Score 15; DB 1;
Pred. No. 1.3e+03;
                                                                                                                      Mismatches
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                                                                                                                                                               The present invention relates to enzymatic nucleic acids which capacitically cleave RNA derived from Hapatitis C virus (HCV). The Expectifically cleave RNA derived from Hapatitis C virus (HCV). The CC enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin CC (HP) motif where the binding arms comprise sequences complementary to one of the substrate sequences defined in the specification. The HCV ribozymes are useful for modulating the expression and/or replication of CC HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV ribozymes are also useful for treating condition associated with HCV infection in conjunction with one or more other drug therapies, particularly type I interferon. Sepecially conterferon alpha, beta or gamma or consensus interferon. The present CC sequence represents a substrate for a HCV hammerhead (HH) ribozyme. Note: Some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was CC obtained in electronic format directly from the USPTO web site at
                                                           Matches
                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therapy; type I interferon; interferon alpha; interferon beta; cytostatic; interferon gamma; consensus interferon; hepatotropic; antiinflammatory; substrate; hammerhead ribozyme; HH ribozyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-MAR-1999;
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                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ROBE/)
(PAVC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MCSW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2002082225-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis C virus substrate #1319 for HCV hammerhead ribozyme #1319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABX03406;
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                                                                                                                                                       seqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MACE/)
                                                           Local Similarity hes 0; Conserv
                            4464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BLATT L.
MCSWIGGEN J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MACEJACK D
                                                                                                                         15
                                                                                                                                                                                                                                                                                                                                                                                                                          Page 64; 80pp; English.
                            TTTTTTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen JA,
                                                                                                                         ₿₽;
                                                           Conservative
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                                                                                                                         0 A;
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                                                                                                                       0 C; 0 G; 0 T;
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                                                          Pred.
15; M
                                                                                        Score
                                                           Mismatches
                                                                                         15;
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                                                      1.3e+03;
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                                                                                                                         U; 0 Other;
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                                                                                        Length 15;
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                                                           Indels
                                                           0
                                                           Gaps
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RESULT 2740 ABL57064 ID ABL5706

ABL57064 standard;

DNA;

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RESULT 2741 ABL57054 유 성

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Query Match
Best Local
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Havens JR,
                                                                                                                                                                                                                                The present sequence is of a hydrazine treated hydrazide precursor phosphoramidite 15-mer, designated oligo 035, which was produced in an example from the invention and which includes a synthetic branching amidite compound. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moieties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL57064;
                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compound for binding macromolecule to substrate surface or conjugation targets, contains phosphorous containing reactive group, hydrazide protecting group and benzene ring, and has predefined formula.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Macromolecule;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Page 44; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-404476/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-2000; 2000WO-US022205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-AUG-2001;
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                                                                                                          Similarity
                                                                                                                                                                          15
                                                                                                                                                                                                                  applications
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                                                                                                                                                                       BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mueller-Ibeler J, Schweitzer M, Onofrey TJ, Greef CH, Wang D;
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US041663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hydrazide; immobilisation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "Diethyl 5-{({2-cyanoethoxy})(diisopropylamino)
phosphanyloxy)methyl)isophthalate, synthetic branchin
amidite"
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/mod_base=
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                                                                                                                                                                       A; 0 C; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cy3 dye"
                                                                                   0,
                                                                                                        Score 15;
Pred. No.
                                                                                                                                                                       ູດ
                                                                                                                                                                       15
                                                                                      Mismatches
                                                                                                                                                                       T; 0
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                                                                                                                              DB 1;
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RESULT 2742
ABL57063
ID ABL5706
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AC ABL5706
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AC ABL5706
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AC ABL5706
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                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                   The present sequence is of a trityl deprotected hydrazide phosphoramidite 15-mer, designated oligo 09, which was produced in an example from the invention. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moieties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hydridisation based assays, DNA chip technology and biosensor
                                                                                                                                                                                                                                                                                                         Compound for binding macromolecule to substrate surface or conjugation targets, contains phosphorous containing reactive group, hydrazide protecting group and benzene ring, and has predefined formula.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hydrazide phosphoramidite oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL57054;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                           Sequence 15 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Macromolecule; hydrazide; immobilisation;
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  22-JUL-2002
                    ABL57063;
                                     ABL57063
                                                                                                                                                                                                                                                                                        Example 2;
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                                                                                                                                                                                                                                                                                                                                                                                            (NANO-) NANOGEN INC
                                                                                                     4464
                                                                                   μ
                                     standard;
                                                                                                    TTTTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                        Page 40; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                               Mueller-Ibeler J, Schweitzer M, Onofrey TJ, Greef CH, Wang D;
                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                               2001WO-US041663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
  (first entry)
                                                                                                                                                          0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "6-((2Cyanoethoxy)(diisopropylamino)
phosphanyloxy)-N'-tritylhexanohydrazide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            *tag=
                                     DNA;
                                                                                                                                100.0%;
                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "phosphoramidite linkage"
                                      15
                                                                                  15
                                      ВP
                                                                                                                       0
                                                                                                                                Score 15;
Pred. No.
                                                                                                                       Mismatches
                                                                                                                                         DB 1;
                                                                                                                                1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09
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                                                                                                                                        Length 15
                                                                                                                       Indels
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                                                                                                                       Gaps
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  22-JUL-2002
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Synthetic.
                                                                                                                                                                                                                                                                                                                                       Hydrazide precursor phosphoramidite oligonucleotide 039.
                                                 modified_base
                                                                                                                                                           modified_base
                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                      Macromolecule;
                                                                                                                                                                                                                                                                                                    hydrazide; immobilisation;
/mod_base=
/note= "3'
                                                                      /mod_base= OTHER
/note= "1,3-Bis-(3',5'-bis(ethyloxycarbonyl) benzyloxy)-5
-((2'-cyanoethyl)(diisopropylamino) phosphanyloxymethyl)-
                                                                                                                                                                              /note=
                                                                                                                                                                                                                                 Location/Qualifiers
                                   tag=
                                                                                                                                          /*tag=
                                                                                                                                                                              "phosphoramidite linkage"
OTHER
Cy3 dye"
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WO200214558-A2

21-FEB-2002.

10-AUG-2001; 2001WO-US041663

11-AUG-2000; 2000WO-US022205

(NANO-) NANOGEN INC

Raddatz S, Havens JR, Mueller-Ibeler J, Schweitzer M, Onofrey TJ, Greef CH, Wang D; Bruecher ņ Windhab z

WPI; 2002-404476/43.

Compound for binding macromolecule to substrate surface or conjugation targets, contains phosphorous containing reactive group, hydrazide protecting group and benzene ring, and has predefined formula.

Example 3; Page 43; 120pp; English

The present sequence is of a hydrazine treated hydrazide precursor phosphoramidite 15-mer, designated oligo O39, which was produced in an example from the invention. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moieties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and biosensor applications

Sequence 15 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;

Query Match Best Local S Matches 15 4464 TTTTTTTTTTTTT 4478 15; بر Similarity Conservative 0.2%; 0 Score 15; DB 1; L Pred. No. 1.3e+03; Mismatches <u>,</u> Length 15; Indels 0, Gaps

0

RESULT 2743 ABL57066 ABL57066; ABL57066 standard; DNA; 15 ВP

(first entry)

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ABL57059
                                                                                                                  RESULT 2744
                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                             The present sequence is of an amino-C6-modified and Cy3 dye labeled T15 oligonucleotide that was used in a comparison of hydrazine and amine attachment moletles on active ester surfaces in an example from the invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acide, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moieties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and biosensor
                                                                                                                                                                                                                                                Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                          Example 12; Page 57; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-404476/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Raddatz S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
Macromolecule; hydrazide; immobilisation; ss
                      Hydrazide precursor phosphoramidite oligonucleotide O33.
                                              22-JUL-2002
                                                                     ABL57059;
                                                                                             ABL57059
                                                                                                                                                                                                                                                                      apprications
                                                                                                                                                                                                                                                                                                                                                                                                                                                  protecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                               targets,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Havens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-AUG-2000; 2000WO-US022205
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Macromolecule; hydrazide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amino-C6-modified and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-AUG-2001; 2001WO-US041663
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                                                                                                                                                                           4464
                                                                                                                                                     -
                                                                                                                                                                                                   l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                              for binding macromolecule to substrate surface or conjugation contains phosphorous containing reactive group, hydrazide ng group and benzene ring, and has predefined formula.
                                                                                          standard;
                                                                                                                                                     TTTTTTTTTTTTT 15
                                                                                                                                                                                                                                                B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mueller-Ibeler J,
Onofrey TJ, Gree
                                                                                                                                                                                                Conservative
                                              (first
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                                                                                                                                                                                                                                                A,
                                                                                          DNA;
                                             entry)
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"Amino-C6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            labeled T15 oligonucleotide
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Cy3 dy
                                                                                                                                                                                                 <u>,</u>
                                                                                                                                                                                                            Score 15;
Pred. No.
                                                                                                                                                                                                                                               15
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                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modification"
                                                                                                                                                                                                                                               T; 0
                                                                                                                                                                                                                                                U; 0 Other;
                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                            1.3e+03
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                                                                                                                                                                                                                       Length 15;
                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                          RESULT 2745
ABL57061
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Best Local S
Matches 15
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         Macromolecule; hydrazide; immobilisation;
                                 Hydrazide
                                                                                                                                                                                                                                                                                                                                                                                                                                                 protecting
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Havens JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Raddatz
                                                        22-JUL-2002
                                                                               ABL57061;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Compound
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                                                                                                                                                                                                            15;
                                                                                                                                                                _
                                 precursor phosphoramidite oligonucleotide
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phosphoramidite 15-mer, designated oligo 033, which was produced in an example from the invention. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moleties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and
                                                                                 ABL57061 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; Page 43; 120pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is of a hydrazine treated hydrazide precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-AUG-2001; 2001WO-US041663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-FEB-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                        4464 TT7
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                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for binding macromolecule to substrate surface or conjugation contains phosphorous containing reactive group, hydrazide ng group and benzene ring, and has predefined formula.
                                                                                                                                                                                                                                                                                                                                                                                                                applications
                                                                                                                                                                                                                                                                                                                                                                       BP; 0 A; 0 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mueller-Ibeler J, Schweitzer M, Onofrey TJ, Greef CH, Wang D;
                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                     TTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000WO-US022205
(first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phosphanyloxymethyl)-benzoic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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/mod_ba
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phosphanyloxymethyl)-benzoic acid methyl es
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note=
entry)
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                                                                                                                                                                                                                                                                                                       0.2%; Score 15;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                     G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                              DB 1;
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                                                                                                                                                                                                                                                                                                       1.3e+03;
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                                                                                                                                                                                                                                                                                                                            Length 15;
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RESULT 2746
ABL57056
ID ABL5705
XX
AC ABL5705
XX
DT 22-JUL-
XX
DE Hydrazi
XX
KW Macromo
XX
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                                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                   The present sequence is of a hydrazine treated hydrazide precursor phosphoramidite 15-mer, designated oligo 037, which was produced in an example from the invention. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moleties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and
                                                                                                                                                                                                                                                                                                                                                                                                                 Raddatz S,
Havens JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Key
modified_base
                                                                                                                                                                                                                                                                                                                                                         Compound for binding macromolecule to substrate surface or conjugation targets, contains phosphorous containing reactive group, hydrazide protecting group and benzene ring, and has predefined formula.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-AUG-2001; 2001WO-US041663
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                                                                                                                                                                                                                                                                                                                                         Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-404476/43.
          Macromolecule; hydrazide; immobilisation; вз
                            Hydrazide phosphoramidite oligonucleotide O31.
                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200214558-A2
                                               22-JUL-2002
                                                                  ABL57056;
                                                                                    ABL57056 standard; DNA; 15 BP
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                                                                                                                                                    4464
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                                                                                                                                                    TTTTTTTTTTTTT 4478
                                                                                                                                                                                                          15 BP;
                                                                                                                                                                                                                             applications
                                                                                                                                                                                                                                                                                                                                         Page 43; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                 Mueller-Ibeler J,
Onofrey TJ, Gree
                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000WO-US022205
                                               (first entry)
                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "1,3-Big-(3',5'-big(ethyloxycarbonyl)
phenylcarbonylamido)-2-((2''-cyanoethyloxy)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diisopropyl)amino)-phosphanyloxy)-propane"
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                                                                                                                                                                              100.0%;
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== "3' Cy3 dye"
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                                                                                                                                                                                         0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            σ
                                                                                                                                                                                                                                                                                                                                                                                                                 er J, Schweitzer M,
Greef CH, Wang D;
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                                                                                                                                                                      0
                                                                                                                                                                     Score 15; DB; Pred. No. 1.3
                                                                                                                                                                                                           G; 15
                                                                                                                                                                                                          T; 0
                                                                                                                                                                                       DB 1;
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                                                                                                                                                                               1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                            Bruecher C,
                                                                                                                                                                                        Length 15;
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RESULT 2747
ABL57060
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Matches
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Havens JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is of a trityl deprotected hydrazide phosphoramidite 15-mer, designated oligo 031, which was produced in an example from the invention. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moisties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and biosensor
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Macromolecule; hydrazide; immobilisation; ss
                                      Hydrazide
                                                                          22-JUL-2002
                                                                                                                     ABL57060
                                                                                                                                                         ABL57060 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                    applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-404476/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-AUG-2000; 2000WO-US022205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-AUG-2001; 2001WO-US041663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-FEB-2002.
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                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Page 40; 120pp; English.
                                      precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mueller-Ibeler J,
Onofrey TJ, Greef
                                                                                                                                                                                                                                                                                                                                                                                                                B₽;
                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                              (first
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/mod_base= OTHER
/note= "3' Cy3 dye"
                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "6-((2Cyanoethoxy)(diisopropylamino)
phosphanyloxy)-N'-tritylhexanohydrazide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1. .15
/*tag=
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                                      phosphoramidite oligonucleotide 034.
                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                            0.2%;
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                15 T;
                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                     1.3e+03;
thes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                          Length 15;
                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or conjugation
, hydrazide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Windhab
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Synthetic

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RESULT 2748
ABK98141
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                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                   The present sequence is of a hydrazine treated hydrazide precursor phosphoramidite 15-mer, designated oligo 034, which was produced in an example from the invention. The invention describes an improved process for immobilisation of macromolecules including DNA, RNA, peptide nucleic acids, pyranosyl-RNA and peptides, especially macromolecules containing multiple reactive sites, to a substrate surface or other conjugation target. It also describes the preparation of oligos containing one or more hydrazides, which can be used for conjugation to surface binding moieties, or for other conjugation reactions. The process is useful e.g. in nucleic acid hybridisation based assays, DNA chip technology and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
modified_base
        gene expression, regulatory sequence; pathogenic double-stranded pathogenic bacteria; virus; replication; virulence; cancer;
                                                                                       07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Compound for binding macromolecule to substrate surface or conjugation targets, contains phosphorous containing reactive group, hydrazide protecting group and benzene ring, and has predefined formula.
                                Triple-helix formation; purine-rich target sequence;
                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-404476/43.
                                                             Triple helix
                                                                                                                  ABK98141;
                                                                                                                                         ABK98141 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3;
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                                                                                                                                                                                                                                    4464
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                                                                                                                                                                                                                         TTTTTTTTTTTTT 4478
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                                                                                                                                                                                                                                                                                                                                           applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 43; 120pp; English
                                                                                                                                                                                                                                                                                                                  B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mueller-Ibeler J, Schweitzer M, Onofrey TJ, Greef CH, Wang D;
                                                                                                                                                                                                                                                              Conservative
                                                           forming
                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Diethyl 5-(((2-cyanoethoxy)(diisopropylamino)
phosphanyloxy)methyl)isophthalate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "3' Cy3 dye"
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                                                                                                                                                                                                                                                                                                                 P.
                                                                                                                                                                                                                                                                                                                 0 C; 0 G; 15 T; 0
                                                           associated oligonucleotide
                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                       0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "phosphoramidite linkage"
                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                          Score 15;
Pred. No.
                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                         DB 1; Le
. 1.3e+03;
                                                                                                                                                                                                                                                                                                                 U; 0 Other;
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                                                                                                                                                                                                                                                                                     Length 15
                                                                                                                                                                                                                                                            Indels
                                   double-helix DNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Windhab
                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                            Gaps
                     DNA;
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                                                                                                                                                                                              RESULT 2749
ABK98184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to methods and oligonucleotides for forming CC a triple-helix comprising a double helical nucleic acid comprising first and second substantially complementary strands, and an oligonucleotide bound to a purine-rich target sequence within the double helical nucleic acid, where the oligonucleotide binds in a parallel and antiparallel corientation, respectively, to target sequences on alternate strands of the double helical nucleic acid. The method has therapeutic applications, where gene expression is controlled by selective triple-helix formation within expression regulatory sequences of a target gene. The coligonucleotides can be used to of a target gene. The coligonucleotides can be used to open triple-helices, and are useful to detect the presence or absence of specific sequences within genomic DNA for diagnostic and therapeutic purposes. The oligonucleotides can be selected to specific sequences required by pathogenic double-stranded DNA including specific sequences required by pathogenic bacteria or viruses for replication or virulence, reducing their pathogenicity. Alternatively, the oligonucleotide can be chosen to target a unique sequence of the
                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pathogen which is not found in the genome of pathogen's host. The oligonuclectides can be used in cancer treatment by way of triple-helix suppression of specific oncogenes including those of endogenous or viral origin. Such therapeutic oligonucleotides are capable of forming triple-helices with such sequences in cancerous cells containing the activated oncogene, so preferentially killing or repressing the cancer causing cell. The present sequence represents an oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5
                                                                    Triple helix
                                                                                                        07-OCT-2002
                                                                                                                                          ABK98184;
                                                                                                                                                                              ABK98184 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Fig 3B; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A triple-helix comprising a double helical nucleic acid (DHNA) and an oligonucleotide which binds in parallel and antiparallel orientation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-536030/57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6403302-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         respectively, for targetting sequences on alternate strands of DHNA to control gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dervan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CALY ) CALIFORNIA INST OF TECHNOLOGY
                                                                                                                                                                                                                                                                                              4464 TTTTTTTTTTTTTT 4478
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                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 0 A; 0 C; 0 G; 15
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beal PA;
                                                                forming associated oligonucleotide #48.
                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-00946976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9305-00168920
                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                              15
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                                                                                                                                                                                                                                                                                                                                                               Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                    T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                    U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                               DB 1; L
1.3e+03;
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                                                                                                                                                                                                                                                                                                                                              Indels
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Triple-helix formation; purine-rich target sequence; double gene expression; regulatory sequence; pathogenic double-str pathogenic bacteria; virus; replication; virulence; cancer;

double-stranded

double-helix DNA, ble-stranded DNA;

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                                                                                                                                                                                            ABZ37501
                                                                                                                                                                                                            RESULT 2750
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                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                    replication or virulence, reducing their pathogenicity. Alternatively, the oligonucleotide can be chosen to target a unique sequence of the pathogen which is not found in the genome of pathogen's host. The oligonucleotides can be used in cancer treatment by way of triple-helix suppression of specific oncogenes including those of endogenous or viral origin. Such therapeutic oligonucleotides are capable of forming triple-helices with such sequences in cancerous cells containing the activated oncogene, so preferentially killing or represent the cancer causing cell. The present sequence represents an oligonucleotide used in the methods of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A triple-helix comprising a double helical nucleic acid (DHNA) and an oligonucleotide which binds in parallel and antiparallel orientation, respectively, for targetting sequences on alternate strands of DHNA to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oncogene suppression; cancerous cell; cytostatic; antimicrobial; ss
                                                     Oligonucleotide SEQ ID NO:622.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-536030/57
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               Library; cleavage; display; diverse family; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 7; Fig 24A; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             control gene expression.
                                                                                                                                                                      ABZ37501 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 BP;
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                                                                                                                                                                                                                                                                                                         4464
                                                                                                                                                                                                                                                                                                                                             l Similarity
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                                                                                                                                                                                                                                                                       TITTTTTTTTTT 15
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                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                         4478
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; Pred. No. 1.3e+(
0; Mismatches
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RESULT 2751
ABV74142/c
ID ABV7414
XX
AC ABV7414
XX
AC ABV7414
XX
AC ABV7414

ABV74142;

ABV74142 standard; DNA; 15

23-JAN-2003

(first entry)

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TITTTTTTTTTTTT 15

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                                                                                                                                                        The present invention describes a method for cleaving single-stranded concleic acid sequences at a desired location. Also described: (1) methods concleic acid sequences at a desired location. Also described: (1) methods concleic acid sequences at a desired location. Also described: (1) methods concleic acid sequence concleid. The family of peptides, collectively displaying at least a part of the diversity of the family, concleic acid that has been cleaved at a concided at least in part by a nucleic acid that has been cleaved at a concider; (3) a method for preparing a library comprising a collection of concider; (3) a method for preparing a library comprising a collection of concider; (3) a method for preparing a library of a diverse family of peptides, concided artibody variable region linked to a version of FIII anchor which does contend for producing a population or a library of immunoglobulin genes; and (6) a library of immunoglobulins that comprise members having at cast one variable domain in which at least one of CDR1 and CDR2 contain conception sequence one of contain sequences that display and/or express a diverse family of peptides or genetic packages that display and/or express a diverse family of contain conception, which can be subsequently used to produce libraries or peptides, polypeptides or proteins. ABZ36912 to ABZ37510 and ABP55464 to invention.
                                        Best Local Similarity Matches 15; Conserv
                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (LADN/) LADNER R C.
(COHE/) COHEN E H.
(NAST/) NASTRI H G.
.(ROOK/) ROOKEY K L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cleaving single-stranded nucleic acid sequences at a desired location contacting the nucleic acid with an single strand oligonucleotide complementary to a nucleic acid region where cleavage is desired.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-APR-2001; 2001US-00837306
24-OCT-2001; 2001US-00000516
25-OCT-2001; 2001US-00045674
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                                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 481; 485pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hoogenboom HRJM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ladner RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HOOG/) HOOGENBOOM H
 4464 TITTITTTTTTT 4478
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                                         Conservative
                                                                                                                BP; 0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                       100.0%;
                                                                                0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nastri HG,
                                         Score 15; DB 1; Le
; Pred. No. 1.3e+03;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rookey KL,
                                                                            Length 15;
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                                           Indels
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                                                                                                                                                                                                  CC primer in PCR, and the common 5' region present in every cDNA clone
CC serves as the 5' priming site. Denaturing and washing leave anchored cDNA
CC in each well. The library is now arrayed and normalised. The method was
CC used to identify and isolate clones encoding G-protein coupled receptors,
CC confectorly pathway of invertebrates and active effectors involved in the
CC olfactory pathway of invertebrates and vertebrates, e.g. odourant binding
CC proteins, or other olfactory or neuronal proteins. The identified
CC receptors and proteins are useful for identifying compounds that reduce a
CC target animal's sensitivity to odours, for manufacturing compounds or
CC devices that mask odours, or trapping invertebrates with odourants.
CC Semiochemicals (e.g. arometics or pheromone mimetics) can be developed
CC with desirable effects on specific species, for the development of pest
CC monitoring systems or non-toxic, species-specific pesticide alternatives,
CC for controlling insect feeding and breeding behaviour, detecting the
CC presence of small air-borne molecules, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is that of the 5' end of a cDNA clone isolated from a cDNA library e.g. a mosquito antenna library. A clone was isolated using a method designed to rapidly array and normalize a complex cDNA library obtained from a target species. Clones are arrayed into multi-
c well plates. Each well contains 16 oligonucleotides (see ABV74137) with a 5' polylinker, a poly-T run capable of binding cDNAs by their poly-A tail and a unique 3' sequence, which allows an anchored oligonucleotide in c each well to selectively hybridise only to those cDNA clones with a c comprehensive level of degeneracy since they are diverse and numerous c enough to ensure that every possible cDNA sequence can be bound by an individual, specific oligonucleotide in a single well. The cDNA library is heated to denature the clones into single stranded DNA, and an aliquot is added to every well. The anchored oligonucleotide serves as the 3'
                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying nucleic acid encoding novel sex-linked-tissue-linked receptors, useful for isolating odorant binding proteins or pesticide alternatives, by analyzing sequences from a male- and female-specific nucleic acid library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G-protein coupled receptor; odourant; receptor; olfaction; array; microarray; anosmia; attractant; arometic; pesticide; ss.
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                                                                                                                                                                 Sequence 15
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31-JAN-2002; 2002US-0353392P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 5; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-029930/02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-2002; 2002WO-US009559
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                                                                                                      Local
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                                                                                   l Similarity
15; Conser
                                         TTTTTTTTTTTT 4478
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                                                                                                                                                                    BP;
                                                                                     Conservative
                                                                                                                                                                    15
                                                                                                                                                                    A; 0 C; 0
                                                                                                  100.0%;
                                                                                0;
                                                                                                      Score 15;
Pred. No.
                                                                                                                                                                 G; 0 T; 0 U; 0 Other;
                                                                                     Mismatches
                                                                                                                          DB 1;
                                                                                                    1.3e+03;
                                                                                                                        Length 15
                                                                                   Indels
                                                                                   0
                                                                                   Gaps
                                                                                   0
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CC The present sequence is that of a poly-T oligonucleotide used in a method CC designed to rapidly array and normalize a complex cDNA library obtained CC from a target species. Clones are arrayed into multi-well plates. Each CC well contains 16 oligonucleotides with a 5' polylinker, a poly-T run CC capable of binding cDNAs by their poly-A tail and a unique 3' sequence, which allows an anchored oligonucleotide in each well to selectively compice 3' key sequences are designed to give a comprehensive level of degeneracy since they are diverse and numerous enough to ensure that every possible cDNA sequence can be bound by an individual, specific CC oligonucleotide in a single well. The cDNA library is heated to denature the common 5' region present in every cDNA clones erves as the 3' primer in PCR, and CC the common 5' region present in every cDNA clone serves as the 5' priming site. Denaturing and washing leave anchored cDNA in each well. The CC library is now arrayed and normalised. The method was used to identify condurant receptors, and active effectors involved in the olfactory or neuronal proteins. The identified receptors and proteins are useful for identifying compounds that reduce a target contents are useful for identifying compounds that reduce a target that mask odours, or trapping invertebrates, e.g. odourant binding proteins, compounds or devices contents as ensitivity to odours, for manufacturing compounds or devices that mask odours, or trapping invertebrates with odourants.

CC semiochemicals (e.g. arometics or pheromone mimetics) can be developed controlling insect feeding and breeding behaviour, detecting the corresence of small air-borne molecules, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABV74141 standard; DNA; 15 BP.
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ABV74141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G-protein coupled receptor; odourant; receptor; olfaction; microarray; anosmia; attractant; arometic; pesticide; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 5; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying nucleic acid encoding novel sex-linked-tissue-linked receptors, useful for isolating odorant binding proteins or pesticide alternatives, by analyzing sequences from a male- and female-specific nucleic acid library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-029930/02
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31-JAN-2002; 2002US-0353392P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide used in cDNA library array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dimitratos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/mod_base=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'note= "5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polylinker"
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Query Match

Sequence 15

BP; 0 A;

0 C; 0 G; 0.2%;

U; 0 Other

Score 15; 15 T; 0

DB 1; Length 15;

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RESULT 2753
ABV75865
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                                 The present invention provides a method for deprotection of an oligonucleotide. This involves reacting a protected oligonucleotide, which is preferably covalently attached to a solid support through a linkage, with a deprotection reagent comprising an active methylene compound and an amine reagent. The process and reagent minimise siderections leading to certain impurities that contaminate synthetic oligonucleotides. The present sequence is a T15 phosphoramidite oligonucleotide having a quencher moiety (0) and minor groove binder (CDPI3) at the 3' end, which was synthesised in an example of the invention. This protected oligonucleotide was treated either with 15% ethanolic ammonia or with 3% diethylmalonate (DEM) dissolved in 15% ethanolic ammonia for 2 hours at 5 degrees C. HPLC analysis showed that deprotection without DEM yielded a complex mixture of products containing only 26.5% of the desired product. When DEM was used, 76.8% of the
                                                                                                                                                                                                                                                                     New oligonucleotide deprotection reagent useful for deprotecting oligonucleotide comprises an active methylene compound and an am
                                                                                                                                                                                                                                                                                                                                                                                                                04-MAR-2002; 2002WO-US006739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide T15-Q-CDPI3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV75865;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         19-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200272864-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Deprotection;
Sequence
                                                                                                                                                                                                                                  Example 2;
                                                                                                                                                                                                                                                                                                           WPI; 2003-046740/04.
                                                                                                                                                                                                                                                                                                                                      Nelson
                                                                                                                                                                                                                                                                                                                                                              (PEKE ) PE
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                         product was obtained
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 15 BP;
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                                                                                                                                                                                                                                                                                                                                                                CORP NY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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0 A;
                                                                                                                                                                                                                               25; 46pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= b
/mod_base= OTHER
/note= "3' Q-CDPI3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mod.
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0 C; 0 G; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "phosphoramidite linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      D)
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T; 0
U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                       and an amine
                                waucts containing
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Query Match 0.2%; Best Local Similarity 100.0%; Matches 15; Conservative (

0

Score 15; DB; Pred. No. 1.3

DB 1; 1 3. 1.3e+03

Length 15; Indels

0

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Gaps

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The invention discloses a hairpin sensor comprising a hairpin loop converse repeat arm and a second inverse repeat arm, and a quenchable conjugated inverse repeat arm and a second inverse repeat arm, and a quenchable conjugated inverse repeat arm of the hairpin loop assembly opposite the complementary probe. Also claimed is a microarray comprising the hairpin complementary probe. Also claimed is a microarray comprising the hairpin complementary probe is bound, directly or indirectly, to a support, a kit conjugated is a microarray comprising the hairpin complementary probe is bound, directly or indirectly, to a support, a kit conjugated is a conductive or semi-conductive, including at least one of the particle is conductive or semi-conductive, including at least one of the cabove hairpin sensor assemblies. The hairpin sensor further comprises a functional group joined to the end of the first inverse repeat arm composite the complementary probe, or first spacer opposite the first inverse repeat arm comprises a ligand positioned content of the first inverse repeat arm and the quenchable fluorescing agent, where the ligand is selected from mercapto, hydroxyl, amino, carboxyl, carbo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 2754
ADA14836/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hairpin sensor; hairpin loop; complementary probe; inverse repeat arm quenchable fluorescing agent; microarray; semiconductor; nanocrystal; rhodamine B-labelled dye; detection; gold support; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hairpin sensor useful for detecting a target nucleotide sequence in sample, comprises a hairpin loop assembly including a complementary and a quenchable fluorescing agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hairpin target sequence, #1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA14836;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-596312/56.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BALL/) BALLINGER C T.
(LOCA/) LOCASCIO M.
(LAND/) LANDRY D P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUN-2001; 2001US-0299460P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUN-2002; 2002US-00176055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-JAN-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Page 11; 16pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /bound_moiety= "Hairpin oligonucleotide #1" /note="Forms a doube-stranded region with the hairpin oligonucleotide shown in example 2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  an example of the invention.
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RESULT 2755
ADB68520
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Best Local Similarity
Matches 15; Conserv
The invention relates to a novel method of inhibiting the expression of one or more genes or RNA transcripts by administering at least one oligomucleotide analogue that includes at least one hydroxyproline nucleic acid (HypNA) monomer to a cell or organism or their extracts. To oligonucleotides of the invention may be used to monitor properties including gene expression, respiration, secretion, signalling, ion-channel activity, cell motility, developmental phenotype and tumour regression. Furthermore, they may be utilised to determine the effects particular genes, as antisense or homologous recombination constructs e.g. for creating animal models of disease and finally, for increasing the activity of some enzymes, such as polymerases. The current sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hairpin sensors. The two or more hairpin sensors include complementary probes that are the same or different and respective quenchable fluorescing agents that are the same or different. The two or more hairpin sensors are arranged in a spatially-defined pattern. The sensor and system are useful for detecting a target nucleotide sequence in a sample. Further, the method involves identifying the target nucleotide sequence by the location of the complementary probe to which the target nucleotide sequence binds. The two or more hairpin sensors include complementary probes or quenchable fluorescing agents, that are different. The sequence presented is the hairpin oligonucleotide target sequence, #1, used in an example of the invention.
                                                                                                                                                                                                                                                                                                                 Example 20; Page 234; 240pp;
                                                                                                                                                                                                                                                                                                                                                                     containing
                                                                                                                                                                                                                                                                                                                                                                                                therapy
                                                                                                                                                                                                                                                                                                                                                                                                                              Method
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tumour regression; single-base mismatch; ss; phosphono-peptide nucleic acid; pPNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hydroxyproline nucleic acid; HypNA; PNA; gene expression; respiration; secretion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single-base mismatch oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB68520 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ACTI-) ACTIVE MOTIF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nydroxyproline nucleic acid; HypNA; PNA; peptide nucleic acid;
gene expression; respiration; secretion; signalling;
ion-channel activity; cell motility; developmental phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4464
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                                                                                                                                                                                                                                                                                                                                                                  of inhibiting expression of genes y and determining effects of genes, ning hydroxyproline nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTTTTTTTTTTT 4478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002US-00072975
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 0 T; 0
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                                                                                                                                                                                                                                                                                                                                                                                     y or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ID 10 DNA.
                                                                                                                                                                                                                                                                                                                                                                                          RNA transcripts, useful y administering oligomers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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Query Match
Best Local
                                                                                   is that of the single-base mismatch oligonucleotide SEQ ID 10 DNA of the invention. This sequence may also comprise a peptide nucleic acid (PNA), a phosphono-PNA (pPNA) or a HypNA.
                                                     Sequence 15
Similarity
                                                     BP; 0
                                                   A; 0 C; 0 G; 15 T; 0
100.0%;
                 0.2%;
Score 15;
Pred. No.
                                                     U; 0 Other;
 1.3e+03
                   DB 1;
                 Length 15;
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RESULT 2756
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Matches

15;

Conservative

0,

Mismatches

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0,

Gaps

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ADC18592
                            nucleic
genomic
                              annealing control primer; ACP; annealing specificity;
nucleic acid amplification; hybridisation; DNA finger
genomic DNA; RNA fingerprint; primer; ss.
Synthetic
                                                                                      Annealing
                                                                                                                      18-DEC-2003
                                                                                                                                                    ADC18592;
                                                                                                                                                                                  ADC18592 standard; DNA;
                                                                                          control
                                                                                                                      (first entry)
                          fingerprint;
                                                                                      primer Oligo-dT15
                                                                                                                                                                                  15
                                                                                                                                                                                  ВÞ
                                                                                          SEQ
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                                            DNA fingerprinting;
                                                                                          NO:54
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Chun 98-DEC-2001; 01-MAY-2002; 19-SEP-2002; 19-JUN-2003. WO2003050305-A1 (SEEG-) SEEGENE INC 2001WO-KR002133. 2002WO-KR000816. 2002WO-KR001781.

Annealing control primer to improve annealing specificity in nucleic amplification, has region complementary to target, arbitrary nucleoti sequence, regulator with universal base/non-discriminatory base analogs. ide

Example 2; SEQ ID NO 54; 190pp; English.

WPI; 2003-627256/59

CC improving the annealing specificity in nucleic acid amplification. (1)
CC has a 3'-end portion with a nucleotide sequence complementary to a site
CC on a template nucleic acid for hybridisation, a 5'-end portion having a
CC pre-selected arbitrary nucleotide sequence, and a regulator portion
CC between the 3' and 5'-end portions, comprising a universal or nonCC discriminatory base analogue, where the regulator portion is capable of
CC regulating an annealing portion of the primer in association with
CC annealing temperature. (1) is useful for improving annealing specificity
CC in nucleic acid amplification. (1) is useful for amplifying a nucleic
CC acid sequence from a DNA or a mixture of nucleic acids, for selectively
CC amplifying a target nucleic acid sequence from a DNA, and for selectively
CC amplifying a target nucleic acid sequence from a MNA, by reverse
CC transcribing the mRNA and performing an amplification reaction using (1).
CC expressed mRNA in two or more nucleic acid semples, by reverse
CC transcribing the mRNA and performing an amplification reaction using (1).
CC is also useful for applily amplifying a target cDNA fragment
CC comprising a cDNA region corresponding to the 3'-end or 5'-end region of
CC an mRNA, for amplifying a population of full-length double-stranded cDNAs
CC complementary to mRNAs, and amplifying 5'-enriched double-stranded cDNAs The present invention describes an annealing control primer (ACP) for

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 RESULT 2757
AAX55055
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   target nucleotide sequence simultaneously using more than one pair of primers in the same reaction, where the primers are derived from (1), for producing a DNA fingerprint of genomic DNA (gDNA), for producing a RNA fingerprint of an mRNA sample, identifying conserved homology segments in a multigene family from an mRNA sample, and for identifying conserved homology segments in a multigene family from gDNA. (1) is also useful for identifying a nucleotide variation in a target nucleic acid, and for mutagenesis in a target nucleic acid. The present sequence represents a primer which is used in the exemplification of the present invention.
The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complementary to mRNA. (I) is also useful for amplifying more than target nucleotide sequence simultaneously using more than one pair.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                       Disclosure; Page 70; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C/EBP-beta antisense oligonucleotide fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX55055 standard; DNA; 16
                                                                                                                                                                                                            vasoconstriction.
                                                                                                                                                                                                                           New antisense oligonucleotides used in
                                                                                                                                                                                                                                                                  WPI; 1999-229400/19.
                                                                                                                                                                                                                                                                                                       Nyce JW;
                                                                                                                                                                                                                                                                                                                                                                                 09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-1999
                                                                                                                                                                                                                                                                                                                                          (UYEC-) UNIV
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                                                                                                                                                                                                                                                                                                                                            EAST CAROLINA
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98US-00093972.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15; DB 1; Li
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                           treatment of, e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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AAX18365
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer
                                          This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta -N3'; or (X)m5'-(gamma)k-delta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine, cytosine or thymine; gamma = thymine; k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-DCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX18365 standard; DNA; 16 BP
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                                                                                                                                                                                                                                               Disclosure;
                                                                                                                                                                                                                                                                                Peptides having at least two PCR.
                                                                                                                                                                                                                                                                                                                               WPI; 1999-183822/16.
                                                                                                                                                                                                                                                                                                                                                                                              18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-FEB-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RT-PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RT-PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                              Page 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the
                                                                                                                                                                                                                                                                                                                                                                SHUZO
                                                                                                                                                                                                                                                                                                                                                                                              97JP-00208312
                                                                                                                                                                                                                                                                                                                                                                                                                              97JP-00208312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA sequence determination; gene sequence analysis;
                                                                                                                                                                                                                                              19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                ႘
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                GLI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                               new nucleotides - useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                               primers in
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Query Match Best Local S Matches 15

1 Similarity
15; Conserv

Conservative

<u>.</u>

Mismatches

Indels

0

Gaps

0

0.2%; Score 15; 100.0%; Pred. No.

DB 1; L 1.4e+03; U; 0 Other;

Length 16;

Sequence 16

₿P;

0 ð

1 C; 1 G;

14 T; 0

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RESULT 2759
AAX18366
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   SEXEXEXEX
                                                                                                                                                RESULT 2760
AAA34502
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                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta -N3'; or (X)m5'-(gamma)k-delta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine or thymine; N = adenine, guanine or thymine; gamma = thymine; k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptides having at least two new nucleotides - useful as primers in \ensuremath{\mathsf{PCR}} .
Human; adenosine receptor; low adenosine antisense oligonucleotide;
phosphorothioate; impaired respiration; inflammation; allergy;
                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 1 A; 0 C; 1 G; 14 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JP11032765-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RT-PCR primer; DNA sequence determination; gene sequence analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RT-PCR primer of the invention SEQ ID 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX18366 standard;
                                              Human adenosine
                                                                           28-JUL-2000
                                                                                                                                  AAA34502 standard; DNA; 16 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 10; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-183822/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUL-1997;
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                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97JP-00208312
                                              receptor related polynucleotide SEQ ID NO:2191.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA; 16
                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                0.2%;
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                                                                                                                                                                                                                                                                                 Score 15; DB 1;
Pred. No. 1.4e+03
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AAF20624 ID AAF2

AAF20624 standard; DNA;

16

문 S

N

16

Matches Query Match Best Local

Similarity

0.2%; Score 15; 100.0%; Pred. No.

DB 1; Length 16; 1.4e+03;

Conservative

0

Mismatches

Indels

<u>,,</u>

Gaps

0

89 15;

acacacacacacaca eceeeeececece

Sequence 16

BP; 0 A; 4 C; 12 G; 0 T; 0 U; 0 Other;

X A X

AAF20624;

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The present invention describes a new composition comprising an antisense considered cold with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, or antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstruction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, control of the adenosine content of the control of decryadenosine which may metastasise to the lungs, including threast and prostate cancer. The reduction of the adenosine content of the crelease of decryadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32311 to AAA32312 represent the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung, disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
uncleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 1815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences given in the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-205971/18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 539; 1343pp; English
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14-MAR-2001

(first entry)

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The present invention describes low adenosine (A) content antisense coligonucleotides the A is replaced by a 'Universal' or alternative base. (C (I) can have respiratory, bronchodilator, antiinflammatory, analysesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with c lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and c chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and c chemokine receptors, adenosine receptors, bradykinin receptors, central nervous system (CNS) and peripheral nervous and non-nervous system or receptors, binding proteins and malignancy associated proteins. The c receptors, binding proteins and malignancy associated proteins. The c antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergy(se) and/or surfactant hypoproduction which are associated with a disease or c condition selected from pulmonary vasoconstriction, allergy(se) and/or allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), hypertension, emphysema, chronic obstructive pulmonary processed (COPD), and/or cancer. AAF18434 to AAF21543 represent human polynucleotide c the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NYEC-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human C/EBP polynucleotide fragment #2191.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAR-2000; 2000WO-US008020
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invention
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Query Match Best Local S Matches 15

1 Similarity 15; Conserv

Conservative

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0.2%;

Score 15; DB; Pred. No. 1.4

DB 1; 1

Length 16 Indels

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0

Sequence 16 BP; 0 A;

4 C; 12 G; 0 T; 0 U; 0 Other;

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RESULT 2762
ABL57075/c
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                                                                                                                                                    The present sequence is that of a perfectly matched target sequence for a complecular beacon comprising an oligonucleotide probe (see ABL57069) covalently attached at the 3' end to fluorescent dye and at the 5' end to can annoparticle. In the native state, the probe forms a hairpin comprising conformation with hybridised termini. The proximity of the fluorophore and quencher (gold nanoparticle) in the molecular beacon results in complementary stretch of the probe to a target sequence, such as the corresponding to the amount of the central complementary stretch of the probe to a target sequence, such as the complementary stretch of the probe to a target sequence, such as the complementary stretch of the probe to a target sequence, such as the complementary stretch of the probe to a target sequence, such as the complementary stretch of the probe to a target sequence, such as the complementary stretch of the extent of which is proportional to the amount of target sequence, the extent of which is proportional to the invention relates generally to the use of metal surface quenchers can be detected. The invention relates generally to the use of metal surface quenchers can be detected. The invention relates generally to the use of metal surface quenchers can be detected. The invention and diagnostic systems
                                                            Best Loc
Matches
                                                                                            Query Match
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30-MAR-2001; 2001US-0280350P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL57075 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sensitively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-404569/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecular beacon; fluorophore; nanoparticle; nucleic acid detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL57075;
                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sensitively detecting proximity changes in a system that utilizes an interacting fluorophore and quencher, for high sensitivity applications, involves utilizing a metal surface as quencher.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dubertret B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYRQ ) UNIV ROCKEFELLER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-AUG-2001; 2001WO-US041941
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecular beacon target sequence.
                                                                             Local
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                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                           B₽;
                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                         30;
                                                                                                                            15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /bound moiety= "Molecular beacon"
/note=""forms double-stranded region with bases 5-21
sequence in ABL57069"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                            ?
                                                                                                                                                                                                                                                                                                                                                                                                         62pp; English.
                                                                           0.2%;
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                                                                                                                           0 C; 1 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                            0,
                                                                               Score 15; DB 1;
Pred. No. 1.4e+
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ABZ96318
                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the cinitiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an cantiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antianflammatory steroid in a subject, for reducing or depleting levels of or reducing sensitivity to adenosine, reducing levels of adenosine antiinflammatory approached the prophylactic or therapeutic or therapeutic respiratory effects of a subject, for reducing levels of adenosine antiinflammatory steroid in a subject, for reducing levels of adenosine approached the prophylactic or therapeutic prophylactic or prophylactic or prophylactic or prophylactic prophylactic or prophylactic p
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                    lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ
                                                                                                                                                                                    Sequence
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23-OCT-2001;
06-FEB-2002;
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                                                                       hydroxyproline nucleic acid; HypNA; PNA; peptide nucleic acid gene expression; respiration; secretion; signalling; ion-channel activity; cell mottlity; developmental phenotype; tumour regression; hybridisation; ss; serine nucleic acid; Se
                                                                                                                                                                                                                                                                                                                                                                   ADB68508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotide used in nonlinear optical technique
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 20-B; 146pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening candidate binding partner(s) for binding to test molecule applying external force field to sample in homogeneous phase, illuminating sample with light beam(s) at fundamental frequencies, a measuring physical properties.
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                                                 phosphono-peptide nucleic acid;
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modified_base
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                                                                        Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
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                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule modulating VEGF receptor(s) gene expression or stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-OCT-1995;
11-JAN-1996;
                                                                                                                                                                                                                                                                                                   KDR; hammerhead ribozyme; hairpin ribozyme;
tumour angiogenesis; psoriasis; rheumatoid
fms-like tyrosine kinase 1; kinase insert
foetal liver kinase 1; ss.
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(RIBO-)
                                                            26-OCT-1995;
11-JAN-1996;
                                                                                                                                                                          01-MAY-1997.
                                                                                                                                                                                                                WO9715662-A2
                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                        Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human flt1 VEGF
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(CHIR )
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2; Conservative
RIBOZYME PHARM INC.
CHIRON CORP.
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96US-00584040.
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96US-00584040
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                      ribozyme; cleavage;
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                                                                                                                                                                                                                                                                                                                              arthritis; ocular disease; domain containing receptor;
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RESULT 2768
AAV37934
ID AAV3793
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AC AAV3793
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Best Local !
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          PCR primers AAV37933-39 are used in the course of the invention. The specification describes a novel protein isolated from leukocytes of patients with 1gA nephropathy. Oligonucleotides based on the DNA sequence encoding this protein are useful as reagents for diagnosing and treating
                                                                                                                                                                                                                    Ishiwata I
Nomura N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGP). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can 1 treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                                 diagnosing
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nephropathy
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KAZUSA DNA RES INS
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                                                                                               Page
                                                                                                                                                                                                                    Nagase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IgA nephropathy; diagnosis; treatment; PCR primer; ss.
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agase T, Sawada
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                                                                                             33; 41pp; Japanese.
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ada S, Takei
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                                                                                                                               DNA encoding nephropathy.
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Pred. No. 1.5e+03;
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RESULT 2770
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AAV49503
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                                                                         AAV49503-V49507 are primers used in the isolation of a human eosinophil cell activator. This protein and antibodies generated from the protein can be used for treatment and discosis of malignant cumours, parasitic infections, allergic inflammation, eosinophilic pneumonia, rapid onset eosinophilia, and autoimmune diseases. DNA can be used for diagnosis, an the antisense DNA in gene therapy of these disorders. The protein can be
                                            Sequence 17
                                                                                                                                                                                     DNA and encoded protein which activates eosinophil cells - of cancer, parasite infection, autoimmune disease and aller
                                                                                                                                                                                                                         WPI; 1998-333261/29
                                                                                                                                                                                                                                                                                                                        05-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                  Eosinophil cell activator; treatment; diagnosis; malignant tumour; parasitic infection; allergic inflammation; eosinophilic pneumonia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 0 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                      Example 1; Page 64;
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                                                                                                                                                                                                                                                                                                                                                                                                                        onset eosinophilia; autoimmune disease; gene therapy; primer;
  15;
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                                           BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
 Conservative
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          0.2%;
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                                                                potential agonists or
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100.0%; Pred. No.
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 ç,
           Score 15;
Pred. No.
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  Mismatches
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            1.5e+03;
                      DB 1;
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                                                                antagonists
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RESULT 2771
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5' (alpha)n-beta -N3'; or (X)m5' (gammal)-Gelta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine or thymine; pamma = thymine; k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 11; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptides
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                                                                                                                                                                                                                                                      AAA30179 standard;
                06-OCT-1998;
                                        06-OCT-1999;
                                                                                                                                                                             PCR primer
                                                                                          WO200020575-A1
                                                                                                                 Synthetic
                                                                                                                                                      Pollenosis-associated protein; high pollen-specific immunoglobulin
                                                                                                                                                                                                      16-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TAKI ) TAKARA SHUZO
                                                                                                                                                                                                                                                                                                                                           4464 TITTTTTTTTTTTT 4478
                                                                                                                                       diagnose;
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                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              having at least two new nucleotides - useful as primers in
                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                                                                             GT15A used in pollenosis associated
                                                                                                                                                                                                                                                                                                                                                                                                                    B₽;
                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                98JP-00284610
                                        99WO-JP005506.
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                                                                                                                                         pollenosis;
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                          treatment;
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                                                                                                                                         human; PCR primer;
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RESULT 2772
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Best Local Sim
Matches 15;
                                                                                                                                   Nagasu T,
Obayashi 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a PCR primer used in the identification of a human pollenosis associated gene. The gene is highly expressed in individuals with high pollen-specific immunoglobulin E (IgE) levels. Invention relates to the nucleotide sequence encoding the pollenosis associated protein, diagnosing pollenosis and screening candidate compounds for treating pollenosis. The gene can be used in diagnosing pollenosis, particularly cedar pollenosis, and screening candidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nagasu T, !
Obayashi I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       compounds
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                                                                                Gene highly expressed in patients with high cedar pollen-specific levels, useful for diagnosing pollenosis, and screening candidate
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                                                                                                                                                                                                                                                                                              IGE;
                                                                                                                                                                                                                                                                                               Pollenosis-associated protein; high pollen-specific immunoglobulin IgE; diagnose; cedar pollenosis; treatment; human; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                             PCR primer GT15C used in pollenosis associated gene identification.
                                                                                                                                                                                                                                                                                                                                                   16-AUG-2000
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                                                                                                                2000-317712/27.
                                                                                                                                                                                                                                                                                              diagnose;
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                                                                      useful for diagno
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Imai Y,
                                                                                                                                    Sugita
., Imai
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                                                 44pp;
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Lu N, Ogawa K;
                                                                                                                                   Kashiwabara T,
Lu N, Ogawa K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4478
                                                                       treatment.
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This sequence represents a PCR primer used in the identification human pollenosis associated gene. The gene is highly expressed in individuals with high pollen-specific immunoglobulin E (19E) leve invention relates to the nucleotide sequence encoding the pollenous transfer of the pollenous transfer in the pollenous transfer of t

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RESULT 2773
AAX82722
      Matches
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Best Local
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Best Local :
                                                                                                                                          This invention describes novel DNA sequences preferentially expressed in IgA nephropathy patients, and DNA sequences stringently hybridizing to them. Independent claims cover diagnostic reagents for IgA nephropathy incorporating the antisense sequences; the treatment of IgA nephropathy using the antisense sequences for mRNA inhibition; proteins associated with IgA nephropathy, containing sequences encoded by the DNA sequences; antibodies recognizing these proteins; the production of the proteins by culture of host cells transformed with DNA encoding them; diagnostic reagents for IgA nephropathy containing the antibodies; and compositions for the treatment of IgA nephropathy which contain the antibodies. The products of the invention can be used for the diagnosis and treatment of IgA nephropathy. This sequence represents a primer used in the isolation and identification of the human IgA nephropathy-associated proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ishiwata 1
Sawada S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    associated protein, diagnosing pollenosis and screening candidate compounds for treating pollenosis. The gene can be used in diagnosing pollenosis, particularly cedar pollenosis, and screening candidate compounds for pollenosis treatment
                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 170; 180pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA sequences preferentially expressed in IgA nephropathy patients, proteins encoded by them, and antibodies to those proteins.
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                                                                                                                                 described in the method of the invention
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human; primer; ss.
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                      Local Similarity
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    15;
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                                                                                      BP; 0 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
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                      0.2%;
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Shibata K, Furuya A;
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  Score 15; DB 1; I
Pred. No. 1.5e+03;
0; Mismatches 0;
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4464 TTTTTTTTTTTTTT 4478

AAZ89372 standard; DNA; 17

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15-JUN-2000 AAZ89372;

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RESULT 2775
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AAX82720
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                                                                                                                                                                                                                         IgA nephropathy patients, and DNA sequences stringently hybridizing to them. Independent claims cover diagnostic reagents for IgA nephropathy incorporating the antisense sequences; the treatment of IgA nephropathy using the antisense sequences for mRNA inhibition; proteins associated with IgA nephropathy, containing sequences encoded by the DNA sequences; antibodies recognizing these proteins; the production of the proteins by culture of host cells transformed with DNA encoding them, diagnostic reagents for IgA nephropathy containing the antibodies; and compositions for the treatment of IgA nephropathy which contain the antibodies. The products of the invention can be used for the diagnosis and treatment of IgA nephropathy. This sequence represents a primer used in the isolation and identification of the human IgA nephropathy-associated proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                   Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA sequences preferentially expressed in IgA nephropathy patients, proteins encoded by them, and antibodies to those proteins.
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Sawada S, Takei M, Shibata K, Furuya A;
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                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel DNA sequences preferentially expressed in
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                                                                                         4464 TTTTTTTTTTTTT 4478
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                                                                                                                                      Score 15;
Pred. No.
                                                                                                                         Mismatches
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                                                                                                                                       1.5e+03;
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XX AAZ3673
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XX Anchore
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KW autoimm
KW prolife
KW prolife
KW growth
XX Synthet
XX W099559
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PN W099559
PN W797-APR-
XX 27-APR-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel method for analysis of an RNA sample which comprises amplifying cDNA with first and second differently labeled primers and analysis of the amplified labeled cDNA. The method is useful for analyzing differential gene expression, for identifying and/or characterizing pharmacological activities or for identifying target genes. The use of different primer combinations allow more cDNAs to be amplified. The method also provides a more detailed analysis than prior arthodom. This sequence represents a primer used to illustrate the
                                                                                                                                                                                                                                                                                                     Stimulus-regulated nucleic acid; sequence profile; nucleic acid level; differentially expressed nucleic acid; disease state; cancer; autoimmune disease; infectious disease; aging; developmental disorder; proliferative disorder; neurological disorder; toxicity; primer; treatment resistance; differential expression; drug discovery; growth factor; epidermal growth factor; radiation; stress; pathogen; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-SEP-1998;
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                          27-AUG-1998;
04-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anchored oligo(dT)
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                                                                        27-APR-1998;
                                                                                                                                                                    04-NOV-1999
                                                                                                                                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                        98US-0083331P.
98US-0098070P.
99US-0118624P.
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two differently labeled primers.
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hes 0;
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Matches
                                                                                     Query Match
                                                                                                                                                      target. The method comprises contacting a probe with an arbitrarily or statistically sampled target and detecting the amount of specific binding of the target to the probe. The methods can be used to identify differentially expressed nucleic acid molecules associated with disease states, such as cancer, autoimmune disease, infectious disease, sidny, developmental disorder, proliferative disorder or neurological disorder. Alternatively the methods can be used to assess the efficacy or toxicity of or a resistance to a treatment. Also the methods can be used to determine differential expression of nucleic acid molecules in response to a stimulus, e.g. a chemical, drug or growth factor (especially epidermal growth factor), radiation, stress or a pathogen. The methods can be used to determine co-regulated genes that can be potential
                                                                                                                                                                                                                                                                                                                                             method
                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 91; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                     acid
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                                                                                                                  Sequence
                                                                                                                                                targets
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Measuring expression
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                                                                       Local
                             4464
                                                                                                                                                                                                                                                                                                                      for measuring the level of two or
The method comprises
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                                                                                                                                                for drug discovery
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                                                                                                                                                                                                                                                                                                                                             represent oligo(dT) primers used for modified differential the method of the invention. The specification describes a measuring the level of two or more nucleic acid molecules i
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                                                          Conservative
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                                                                       Score 15; DB 1;
Pred. No. 1.5e+0
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                                                                        1.5e+03
                                                                                      Length 17;
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26-APR-2000;

2000WO-JP002730

27-APR-1999;

99JP-00120489

(GENO-)

GENOX RES

INC

Nagasu T, S Obayashi I,

Sugita , Imai

Kashiwabara T, Os Yoshida N, Ogawa

Oshida T,

۳,

Mateui

Obayashi M,

Gunji

s

2000-687339/67.

Pollinosis-associated gene 373 undergoing significantly low expression subjects with high cedar pollen-specific immunoglobulin-E levels, usefi in diagnosis of allergic diseases and screening drug candidates.

Example 6; Page 69; 80pp; Japanese

The invention relates to the human pollinosis-associated gene 373 which CC exhibits significantly reduced expression in the T-cells of individuals CC with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene CC with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene CC with high cedar pollen from individuals allergic to cedar pollen using CC the differential display method. The invention also relates also relates to the protein encoded by pollinosis gene 373; expression constructs and CC host cells comprising pollinosis-associated gene 373 nucleic acids; pollinosis-associated gene 373 primers and probes; antibodies against the CC protein encoded by the gene; methods of detection of pollinosis of allergic diseases via the detection of pollinosis-associated gene 373 nucleic acids. The invention additionally encompasses methods of screening drug CC candidates for the treatment of allergic disease by measuring the CC expression of pollinosis-associated gene 373 in pollen antigen-stimulated CC T-cells in the presence of a test compound relative to a control. CC diseases and in the screening of drug candidates for the treatment of Such diseases. The present sequence represents a PCR primer used in the control final processor of the present sequence area and primer used in the control control control diseases. The present sequence represents a PCR primer used in the control control control control control control control diseases. of human pollinosis-associated gene 373 cDNA

XXXDDDDDDDDDDDDDDDDDDDXXX

Sequence 17 BP; 1 ۶ 0 C; 1 G; 15 T; 0 u; o Other;

Query Match Best Local S Matches 15 l Similarity Conservative (0.2%; 0 Score 15; DB; Pred. No. 1.5 DB 1; 1.5e+03; 0 Length 17; Indels 0; Gaps 0

밁 RESULT 2778 AAC64203 N S

4464

TTTTTTTTTTTT 4478

16

AAC64203 standard; DNA; 17

AAC64203;

21-FEB-2001 (first entry)

PCR anchor primer, SEQ ID NO:4, used in human gene 373 isolation

Human; pollinosis-associated gene 373; IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced expression; detection; drug screening; allergic disease; PCR primer; ss. detection; diagnosis;

WO200065046-A1

02-NOV-2000

26-APR-2000; 2000WO-JP002730.

27-APR-1999; 99JP-00120489

(GENO-) GENOX RES INC

Obayashi I, Nagasu T, Sugita Y, Kashiwabara Yoshida N, T, Oshida T, Obayashi M, Ogawa K, Matsui K; Gunji

WPI; 2000-687339/67.

Pollinosis-associated gene 373 undergoing significantly low expression i subjects with high cedar pollen-specific immunoglobulin-E levels, useful in diagnosis of allergic diseases and screening drug candidates.

80pp; Japanese

The invention relates to the human pollinosis-associated gene 373 which exhibits significantly reduced expression in the T-cells of individuals

CC was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates also relates constructs and constructs are constructed by the gene; methods of detection of pollinosis allergic considered gene 373 nucleic constitution additionally encompasses methods of screening drug candidates for the treatment of allergic disease by measuring the construction of pollinosis-associated gene 373 in pollen antigen-stimulated construction constructs and in the screening of drug candidates for the treatment of construction constructs and in the screening of drug candidates for the treatment of construction constructs and in the screening constructs and in the screening constructs and constructs and in the screening constructs and construct

Sequence 17 BP; 0 A; 1 C; 1 G; 15 T; 0 U; 0 Other;

Matches Query Match Local 15; Similarity Conservative 0.2%; 0 Score 15; Pred. No. Mismatches DB 1; 1.5e+03; 0, Length 17; Indels ٥, Gaps 0

4464 TTTTTTTTTTTTTT N 16 4478

밁 Ś

RESULT 2779

AAC64181 standard; DNA; 17

AAC64181;

21-FEB-2001 (first entry)

PCR anchor primer, SEQ ID NO:2, used in human gene 419 isolation.

allergic disease; Human; pollinosis-associated gene 419; FAF-1 homologue; Fas-associated factor-1; Ig-cell; reduced expression; factor-1; IgE; immunoglobulin
d expression; detection; diagno primer; ss. diagnosis; E; cedar pollen allergy; drug

Synthetic

WO200065045-A1

02-NOV-2000.

26-APR-2000; 2000WO-JP002729

27-APR-1999; 99JP-00120490

(GENO-) GENOX RES INC.

Nagasu T, S Obayashi I, Sugita Y, Kashiwabara Yoshida N, T, Oshida T, Obayashi M, Ogawa K, Matsui K; Gunji S;

WPI; 2000-687338/67.

Pollinosis-associated gene 419 undergoing significantly low subjects with high cedar pollen-specific IgE levels, useful of allergic diseases and screening drug candidates. expression in in diagnosis

Example 6; Page 49; 77pp; Japanese

The invention relates to the human pollinosis-associated gene 419 which exhibits reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. Pollinosis-associated gene 419 has homology with the gene encoding human Fas-associated

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RESULT 2780
AAC64182
ID AAC6418
XX AAC6418
XX PCR anc
XX PCR anc
XX PCR anc
XX Human;
KW Fas-ass
KW T-cell;
KW allergi
XX Synthet
XX AC10006
XX O2-NOV-
XX PD 02-NOV-
XX PP 27-APR-
XX (GENO-)
XX Nagasu
PI Obayasl
XX Nagasu
PI Obayasl
XX POllin
PT Subject
PT Pollin
PT subject
PT Subject
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The invention relates to the human pollinosis-associated gene 419 which exhibits reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. Pollinosis-associated gene 419 has homology with the gene encoding human Fas-associated factor-1 (FAF-1). The invention also relates to the protein encoded by pollinosis gene 419; expression constructs and host cells comprising pollinosis-associated gene 419 primers and probes; nucleic acids; pollinosis-associated gene 419 primers and probes;
                                                                                                                                                                                                                                                                                                                                   Nagasu T,
Obayashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; pollinosis-associated gene 419; FAF-1 homologue; Fas-associated factor-1; IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced expression; detection; diagnosis; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC64182 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                       Pollinosis-associated gene 419 undergoing significantly low subjects with high cedar pollen-specific IgE levels, useful of allergic diseases and screening drug candidates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-APR-2000; 2000WO-JP002729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             allergic
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                                                                                                                                                                                     Example 6; Page 49; 77pp;
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Yoshida N,
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Ogawa K, Mat
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                                                                                                                                                                                                                                            expression in in diagnosis
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antibodies against the protein encoded by the gene; methods of detection of pollinosis-associated gene 419 nucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 419 nucleic acids. The invention additionally encompasses methods of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 419 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control.pollinosis-associated gene 419 is useful in the diagnosis of allergic diseases and in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-associated gene 419 cDNA
Sequence 17 BP; 0 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
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Query Match
Best Local S
Matches 15
        4464 TTTTTTTTTTTTTTTT
N
                             15;
                                     Similarity
                             Conservative
                                     100.0%;
                                            0.2%;
              4478
16
                              0
                                      Score 15;
Pred. No.
                              Mismatches
                                     DB 1; Lo
                                            Length 17;
                              0;
                             Gaps
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吊 S

RESULT PCR anchor 21-FEB-2001 AAC64171; AAC64171 standard; DNA; 17 primer, (first entry SEQ ä NO:2, used in human ВP gene 513 isolation.

Human; pollinosis-associated gene 513; IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced expression; detection; diagnosis; drug screening; allergic disease; PCR primer; 88.

WO200065049-A1

02-NOV-2000.

27-APR-1999; 99JP-00120491

26-APR-2000; 2000WO-JP002733

(GENO-) GENOX RES INC

Nagasu T, Obayashi : Η Sugita Y, Imai Y, Kashiwabara T, Or Yoshida N, Ogawa Oshida T, <u>~</u> a T, Obayashi M, Matsui K; S

Pollinosis-associated subjects with high ced of allergic diseases a cedar es and gene 513 undergoing significantly low expression dar pollen-specific IgE levels, useful in diagnosiand screening drug candidates. xpression in 1 diagnosis

Example 6. Page 38; 46pp; Japanese

The invention relates to the human pollinosis-associated gene 513 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates to methods of detection of pollinosis-associated gene 513 nucleic acids; a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 513 nucleic acids; and methods of screening drug candidates for the reatment of allergic disease by measuring the expression of pollinosis-associated gene 513 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis-associated gene 513 in pollen antigen-stimulated T-cells in the screening of useful in the diagnosis of allergic diseases and in the screening of trug candidates for the treatment of such diseases. The present sequence

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AAC64172
                                                                             The invention relates to the human pollinosis-associated gene 513 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates to methods of detection of pollinosis-associated gene 513 nucleic acids; and methods of screening drug candidates for the treatment of allergic diseases via the detection of pollinosis-associated gene 513 nucleic acids; and methods of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 513 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis-associated gene 513 in the diagnosis of allergic diseases and in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-
              Query Match
Best Local :
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Matches 15; Conserv
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Obayashi
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Imai Y,
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Conservative
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             0.2%; Score 15;
100.0%; Pred. No.
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Yoshida N,
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Pred. No.
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                                                       T;
                                                      U; 0 Other;
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                           DB 1;
             1.5e+03;
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                          Length 17;
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sui K;
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                                                                                                                                                                                                                       The invention relates to the human pollinosis-associated gene 581 which ce exhibite significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene ce was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates also relates to the protein encoded by pollinosis-associated gene 581; to expression constructs and host cells comprising pollinosis-associated gene 581 mucleic acids; pollinosis-associated gene; methods of detection antibodies against the protein encoded by the gene; methods of detection control of allergic diseases via the detection of pollinosis-associated gene 581 nucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 581 nucleic acids. The invention additionally encompasses methods of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 581 in pollen antigen-stimulated T-cells in the presence of a test compound relative to allergic disease and in the greening of drug candidates for the diagnosis of allergic disease of allergic disease of the detection of pollinosis-associated gene 581 in the diagnosis of allergic disease of the disease of allergic disease of the disease of 
                                                                Query Match
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                                                                                                                                                               allergic diseases and in the screening of drug candidates for treatment of such diseases. The present sequence represents a used in the isolation of human pollinosis-associated gene 581
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pollenosis-associated gene 581 undergoing significantly low subjects with high cedar pollen-specific IgE levels, useful of allergic diseases and screening drug candidates.
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Obayashi I,
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                                                              100.0%;
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100.0%; Pred. No.
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hes 0;
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Matsui K;
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RESULT 2784
AAC64162
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AAC64213
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AXU
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AAC64213;
                         AAC64213 standard; DNA; 17
                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENO-) GENOX RES INC
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                                                                                                                 4464
                                                                                          N
                                                                                                                                                        Similarity
                                                                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sugita Y,
                                                                                                                                                                                              B₽;
                                                                                                                                            Conservative
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                                                                                                                  TTTTTTTTTT 4478
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                                                                                                                                                                                              0 A;
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                                                                                                                                                       0.2%;
                                                                                                                                                                                              1 C; 1 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kashiwabara T, Oshida T,
Yoshida N, Ogawa K, Mat
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                                                                                                                                           0
                                                                                                                                                       Score 15; DB 1; 1
Pred. No. 1.5e+03
                                                                                                                                                                                              15
                                                                                                                                             Mismatches
                                                                                                                                                                                              H;
                                                                                                                                                                                              U; 0 Other;
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                                                                                                                                            0
                                                                                                                                                                   Length 17;
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ui K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression in in diagnosis
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                                                                                                                                           Gaps
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grug
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                                 Pollinosis-associated gene 627 undergoing significantly low expression in subjects with high cedar pollen-specific IgE levels, useful in diagnosis of allergic diseases and screening drug candidates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-FEB-2001
 Example
                                                                                                                WPI; 2000-687344/67.
                                                                                                                                                   Nagasu T,
Obayashi I,
                                                                                                                                                                                                          (GENO-) GENOX RES INC
                                                                                                                                                                                                                                               27-APR-1999;
                                                                                                                                                                                                                                                                                     26-APR-2000; 2000WO-JP002735
                                                                                                                                                                                                                                                                                                                          02-NOV-2000.
                                                                                                                                                                                                                                                                                                                                                               WO200065051-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR anchor primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                      screening;
6; Page 41; 51pp; Japanese.
                                                                                                                                                   Sugita Y,
Imai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                 99JP-00120493
                                                                                                                                                                                                                                                                                                                                                                                                                                          allergic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID
                                                                                                                                                   Kashiwabara T, Oshida T,
Yoshida N, Ogawa K, Mate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO:2, used in
                                                                                                                                                                                                                                                                                                                                                                                                                                        expression; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human
                                                                                                                                                       Mateui
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                                                                                                                                                     Obayashi M,
ui K;
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The invention relates to the human pollinosis-associated gene 627 which CC exhibits significantly reduced expression in the T-cells of individuals CC with high cedar pollen-specific IgE (Ammunoglobulin E) levels. The gene CC was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates to methods of CC diagnosis of allergic diseases via the detection of pollinosis-associated gene 627 nucleic acids; a method of CC gene 627 nucleic acids; and a method of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 627 in pollien antigen-stimulated T-cells in the presence CC associated gene 627 in pollien antigen-stimulated T-cells in the presence CC is useful in the diagnosis of allergic diseases and in the screening of CC drug candidates for the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-

Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;

Query Match Best Local S Matches 15 15; Similarity Conservative 100.0%; 0.2%; 0 Score 15; DB 1; Li Pred. No. 1.5e+03; Mismatches 0 Length 17; 0 Gaps

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       4464 TTTTTTTTTTTTT 4478
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                                                            RESULT 2786
                                21-FEB-2001
                                         AAC64214;
                                                   AAC64214 standard; DNA; 17
                                (first
                                entry)
                                                   ВP
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PCR anchor primer, SEQ ID NO:3, used in human gene 627 isolation

Human; pollinosis-associated gene 627; IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced expression; detection; drug screening; allergic disease; PCR primer; 88.

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                                                                                                                                                                                                                                                                                                                      RESULT 2787
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                           Human; pollinosis-associated gene 795; vimentin homologue; IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced express detection; diagnosis; drug screening; allergic disease; PCR pri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pollinosis-associated gene 627 undergoing significantly low subjects with high cedar pollen-specific IgE levels, useful of allergic diseases and screening drug candidates.
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27-APR-1999;
                                                         02-NOV-2000
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                                                                                                                                                                                                                                                                                          AAC64231 standard;
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Obayashi I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-APR-2000; 2000WO-JP002735
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                           26-APR-2000; 2000WO-JP002734.
                                                                                    WO200065050-A1
                                                                                                                   Synthetic
                                                                                                                                                                                                     PCR anchor primer,
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                                                                                                                                                                                                                                                                                                                                                                                                                          15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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, Imai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                         larity 100.0%; 1 Conservative 0;
                                                                                                                                                                                                                                (first entry)
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99JP-00120494
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                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                   SEQ ID NO:3, used in human gene 795 isolation.
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Yoshida N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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Matsui K;
                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                                                                                           expression;
PCR primer;
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The invention relates to the human pollinosis-associated gene 795 which certainly reduced expression in the T-cells of individuals calibrated from T-cells from individuals allergic to cedar pollen-specific IgE (immunoglobulin E) levels. The gene curve was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. Pollinosis-associated gene 795 has the homology with the human vimentin gene. The invention also relates also relates to the protein encoded by pollinosis-associated gene 795 to expression constructs and host cells comprising pollinosis-associated gene 795 mucleic acids; pollinosis-associated gene 795 rimers and probes; concleic acids; pollinosis-associated gene 795 mucleic acids; pollinosis-associated gene 795 mucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 795 mucleic acids. The invention additionally encompasses methods of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 795 in pollen cantigen-stimulated T-cells in the presence of a latergic disease by candidates for the treatment of allergic disease so do in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-associated gene 795 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nagasu T, Sı
Obayashi I,
Yokoi A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pollinosis-associated gene 795 undergoing significantly low subjects with high cedar pollen-specific IgE levels, useful of allergic diseases and screening drug candidates.
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EISAI
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, Imai Y,
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Yoshida N, Ogawa K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a T, Obayashi M, Gunji
Matsui K, Takahashi E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression in in diagnosis
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Query Match Best Local S Matches 15 4464 15; N Similarity TTTTTTTTTTTTT 4478 Conservative 100.0%; 0.2%; 0; Score 15; DB 1; Pred. No. 1.5e+C 1.5e+03; thes 0; Length 17; Indels 0 Gaps

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Sequence 17

BP; 0 A; 1 C; 1 G;

15 T; 0

U; 0 Other;

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RESULT 2788
AAC64230
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                 AAC64230 standard; DNA; 17
AAC64230
                  ВP
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PCR anchor primer, SEQ ID NO:2, used in human gene 795 isolation 21-FEB-2001

(first entry)

detection; Human; iuman; pollinosis-associated gene 795; vimentin homologue; IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced express letection; diagnosis; drug screening; allergic disease; PCR pri 88.

Synthetic

WO200065050-A1

02-NOV-2000

26-APR-2000; 2000WO-JP002734

27-APR-1999; 99JP-00120494

(GENO-) GENOX RES INC

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RESULT 2789
AAC9229
ID AAC9229
XX AAC9229
XX AAC9229
XX Human p
XX Human p
XX Human p
XX Human p
XX Homo sa
XX AT HOMO sa
XX 18-MAY-
XX 18-MAY-
PF 18-MAY-
XX (GENO-)
PA (GENO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to the human pollinosis-associated gene 795 which CC exhibits significantly reduced expression in the T-cells of individuals CC with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene CC was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. Pollinosis-associated gene 795 has CC homology with the human vimentin gene. The invention also relates also CC constructs and host cells comprising pollinosis-associated gene 795 councleic acids; pollinosis-associated gene 795 primers and probes; CC antibodies against the protein encoded by the gene; methods of detection CC of pollinosis-associated gene 795 primers and probes; CC diagnosis of allergic diseases via the detection of pollinosis-associated gene 795 nucleic acids. The invention additionally encompasses methods of CC gene 795 nucleic acids. The invention additionally encompasses methods of CC exceening drug candidates for the treatment of allergic disease by a control. Pollinosis-associated gene 795 in pollen CC antigen-stimulated T-cells in the presence of a test compound relative to CC acontrol. Pollinosis-associated gene 795 in useful in the dangnosis of allergic diseases and in the screening of drug candidates for the creatment of such diseases. The present sequence represents a PCR primer CC used in the isolation of human pollinosis-associated gene 795 cDNA
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Best Local
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Obayashi I,
  Nagasu T,
                                              (GENO-)
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                                                                                                                                                                  18-MAY-2000; 2000WO-JP003191:
                                                                                                                                                                                                                                                                                                                                                        Human; pollinosis-associated gene 465; pollen scattering; allergy;
allergic disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                             Human pollinosis-associated gene 465 related PCR primer SEQ ID NO:2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC92292,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC92292 standard;
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, Imai Y,
  Sugita Y,
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                                                RES INC.
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                                                                                                                       99JP-00148784
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    Kashiwabara
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wa K, Matsui K, Takahashi E;
    Oshida
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    H
    Obayashi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression in in diagnosis
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    Gunji
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Matches
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Best Local
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Yokoi A;
                                                                                                                                                                                                       The present invention describes the human pollinosis-associated gene 465 which has a nucleic acid sequence of 3442 base pairs (bp), given in (AAC92291), that undergoes significantly low expression in subjects after pollen scattering, and is useful in the diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen. The gene is useful in the diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen. The present sequence represents a PCR primer which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diseases cells to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pollinosis-associated gene 465 undergoing significantly low expression subjects after pollen scattering, useful in diagnosis of allergic diseases and screening candidate compounds to regulate response of T
                                                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                      4464 TTTTTTT
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                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and screening can antigen stimulus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 43;
                                                                                                                                                                     BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Imai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     61pp; Japanese.
                                                                                                        100.0%;
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                                                                                                                            0.2%;
                                      4478
16
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                                                                                                        Score 15;
Pred. No.
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                                                                                      Mismatches
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                                                                                                     DB 1; Le
1.5e+03;
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                                                                                                                          Length 17;
                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                             Nagasu T,
Obayashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                             Pollinosis-associated gene 465 und
subjects after pollen scattering,
diseases and screening candidate of
cells to antigen stimulus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; pollinosis-associated gene 465; allergic disease; PCR primer: as
                                                                                                                                                                                                                                                                       WPI; 2001-061528/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-MAY-2000; 2000WO-JP003191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human pollinosis-associated gene 465 related PCR primer SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC92293 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                            COKOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENOX RES INC
EISAI CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                Sugita Y,
Imai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99JP-00148784
                                                                                                                                                                                                                                                                                                                                                                             Kashiwabara T, Oshio
Yoshida N, Ogawa K,
                                                                                                                                                                  undergoing significantly low expression ng, useful in diagnosis of allergic
                                                                                                                                                                                                                                                                                                                                                                                                                 Oshida T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pollen
                                                                                                                                                                                                                                                                                                                                                                                Matsui K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               scattering; allergy;
                                                                                                                                                                                                                                                                                                                                                                                Obayashi M, Gu
ui K, Takahashi
                                                                                                                                                                                                                                                                                                                                                                                Gunji S;
shi E;
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Example

6; Page 44;

61pp; Japanese

compounds

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response

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RESULT 2791
AAC91720
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Best Local S
Matches 15
Nagasu T, s
Obayashi I,
Yokoi A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes the human pollinosis-associated gene 465 which has a nucleic acid sequence of 3442 base pairs (bp), given in (AAC92291), that undergoes significantly low expression in subjects after pollen scattering, and is useful in the diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen. The gene is useful in the diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen. The present sequence represents a PCR primer which is used in an example from the present invention
                                                                                                          27-MAY-1999;
                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                       PCR anchor
                                                                                                                                                                                                                                                                                                                                                     27-MAR-2001
                                                              (GENO-)
                                                                                                                                        18-MAY-2000; 2000WO-JP003192
                                                                                                                                                                                                   WO200073440-A1.
                                                                                                                                                                                                                                                             allergic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15;
                                                                                                                                                                                                                                                                                       pollinosis-associated gene 787;
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                                                              GENOX
EISAI
                                                                                                                                                                                                                                                                            expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17
                                                                                                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTTTTTTTTTTT
               Sugita Y,
Imai Y,
                                                                                                                                                                                                                                                                                                                     primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                              CO LTD.
                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                          99JP-00148785
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                                                                                                                                                                                                                                                           PCR
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                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                       SEQ ID
                                                                                                                                                                                                                                                                         detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 C; 1 G;
              Kashiwabara
Yoshida N,
                                                                                                                                                                                                                                                             primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
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Pred. No.
                                                                                                                                                                                                                                                             88
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                                                                                                                                                                                                                                                                         diagnosis; drug screening;
                                                                                                                                                                                                                                                                                                                     used in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T; 0
               T, Oshida T,
Ogawa K, Matsu
                                                                                                                                                                                                                                                                                         pollen allergy; T-cell,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                 Mateui
                                                                                                                                                                                                                                                                                                                     gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 17
               Obayashi M,
ui K, Takaha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                       787 isolation.
                 Takahashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
               Gunji
shi E;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                    Nagasu T,
Obayashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           associated gene 787 in pollen antigen-stimulated T-cells in the prese of a test compound relative to a control. Pollinosis- associated gene is useful in the diagnosis of allergic diseases and in the screening drug candidates for the treatment of such diseases. The present seque represents a PCR primer used in the isolation of human pollinosis-associated gene 787 cDNA
                                                                                                                                                                             WPI; 2001-032159/04
                                                                                                                                                                                                                          (GENO-)
                                                                                                                                                                                                                                                                 18-MAY-2000; 2000WO-JP003192.
                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                           PCR anchor primer,
                                                                                                                                                                                                                                                                                                                                                                                           AAC91719;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17
                                                                                                                                                                                             Yokoi
                                                                                                                                                                                                                                                  27-MAY-1999;
                                                                                                                                                                                                                                                                                07-DEC-2000
                                                                                                                                                                                                                                                                                                WO200073440-A1
                                                                                                                                                                                                                                                                                                                             allergic
                                                                                                                                                                                                                                                                                                                                     reduced
                                                                                                                                                                                                                                                                                                                                             Human;
                                                                                                                                                                                                                                                                                                                                                                           27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                          AAC91719 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                4464 TITTTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                                                                    pollinosis-associated gene 787; pollen allergy; T-cell;
d expression; detection; diagnosis; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15;
                                                                                                                                                                                                                          GENOX
EISAI
                                                                                                                                                                                                                                                                                                                             expression;
disease; Po
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                    Sugita
, Imai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP;
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                                                                                                                                                                                                                          CO LTD.
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                                                                                                                                                                                                                                                                                                                                                            SEQ
                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                           entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 C; 1 G;
                                                                                                                                                                                                    Kashiwabara
Yoshida N,
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                                                                                                                                                                                                                                                                                                                                                                                                          17
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                                                                                                                                                                                                                                                                                                                                                           NO:2, used in human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15
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                                                                                                                                                                                                    ı T, Oshid
Ogawa K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Η,
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                                                                                                                                                                                                             Oshida T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                    Matsui
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other,
                                                                                                                                                                                                                                                                                                                                                           gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                    Obayashi M, Gunji
ui K, Takahashi E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                             787
                                                                                                                                                                                                                                                                                                                                                             isolation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence
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d gene 787
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The invention relates to the human pollinosis-associated gene 787 which exhibits significantly reduced expression in the T-cells of individuals after the pollen-scattering season, relative to expression levels in T-cells before the pollen-scattering season. The gene was isolated from T-cells from individuals allergic to pollen using the differential display method. The invention also relates to pollinosis-associated gene 787 primers and probes; methods of detection of pollinosis-associated gene 787 nucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis associated gene 787 nucleic acids. The invention additionally encompasses a method of screening drug candidates for the diseases Pollinosis-associated gene 787 undergoing significantly low expression subjects after pollen scattering, useful in diagnosis of allergic diseases and screening candidate compounds to regulate response of T Example 6; antigen stimulus. Page 40; 54pp; Japanese

expression

WPI; 2001-032159/04.

cells before the pollen-scattering season. The gene was isolated from T-cells before the pollen-scattering season. The gene was isolated from T-cells from individuals allergic to pollen using the differential display method. The invention also relates to pollinosis-associated gene 787 primers and probes; methods of detection of pollinosis-associated gene 787 nucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 787 nucleic acids. The invention additionally encompasses a method of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 787 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis- associated gene 787 rollinosis-associated gene 787 undergoing significantly low expression subjects after pollen scattering, useful in diagnosis of allergic diseases and screening candidate compounds to regulate response of T cells to article. of a test compound relative to a control. Pollinosis- associated is useful in the diagnosis of allergic diseases and in the screen The invention relates to the human pollinosis-associated gene 787 which exhibits significantly reduced expression in the T-cells of individuals after the pollen-scattering season, relative to expression levels in T-Example cells 6; Page 40; antigen stimulus 54pp; Japanese. s- associated gene in the screening o 9 ij

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RESULT 2793
AAC82875
ID AAC8287
XX AAC8287
AC AAC8287
XX Polling
XX PR 27-MAY-
XX (GENO-)
XX Nagasu
PI Obayasi
XX WPI; 20
XX Polling
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AXE
                                                          RESULT 2794
AAC82874
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                                                                                                                                                                                                                           Query Match
Best Local S
Matches 15
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        represents a PCR associated gene
                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel nucleic acid molecule comprising sequence (I) which undergoes significantly low expression in subjuster pollen scattering, and is useful in diagnosis of allergic diand screening candidate compounds for remedies capable of regulativesponse of T cells to the stimulus by an antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pollinosia-associated gene 441 which undergoes lower expression subjects after pollen scattering, useful in diagnosis of allergidiseases and screening candidate compounds to regulate response cells to antigen stimulus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC82875 standard;
  AAC82874;
                                    AAC82874
                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENO-) GENOX RES INC
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                                                                                                                                                                                      4464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 seis; pollinosis-associated gene 441; allergy; T cell;
scattering; antigen; primer; ss.
                                                                                                                                                                                                                           l Similarity
15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
                                                                                                                                             N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Page 35; 42pp; Japanese.
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                                                                                                                                                                                                                                                                                                                17 BP; 0
                                         standard;
                                                                                                                                               TTTTTTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sugita Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 1 A;
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                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA; 17
                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 C; 1 G;
                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                1 C; 1 G; 15 T; 0
                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kashiwabara T,
Yoshida N, Og
                                           17
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Ogawa K, Mat
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Obayashi I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel nucleic acid molecule comprising a sequence (I) which undergoes significantly low expression in subjects after pollen scattering, and is useful in diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pollinosis-associated gene 441 which undergoes lower expression in subjects after pollen scattering, useful in diagnosis of allergic diseases and screening candidate compounds to regulate response of cells to antigen stimulus.
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                                                                                                                                                                                                                                                                                                                                     Nucleotide sequence of primer GT15C.
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, Imai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 35;
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Yoshida N,
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention provides a method of diagnosis of allergies that assaying the levels of expression of genes B1001, B1466, B1072 in T-cells; and comparing them with the level of expression in cells. The method is useful for diagnosing allergies, particula dermatitis. The present sequence represents a PCR primer used f
                                                                                                                                                  Diagnosis
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                                                    ng the levels of expression of genes B1001, B1466, B1072 or B1151 rells; and comparing them with the level of expression in healthy T-The method is useful for diagnosing allergies, particularly atopic itis. The present sequence represents a PCR primer used for its of the expression of the above genes
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                                                                                                          a method of diagnosis of allergies that
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The present invention describes a human genome-derived myosin-like CC protein 1 (hiDMLP-1). The protein and polynucleotide sequences of hdDMLP-CC 1 can be used in gene therapy and vaccine production. The hdDMLP-1 cC nucleic acids can be used as probes to detect, characterise and quantify CC hdDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hdDMLP-1 corotein variants having desired phenotypic improvements, and for cexpressing the proteins. The hdDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hdDMLP cused as immunogens to raise antibodies that specifically recognise hdDMLP cand/or amount specifically of hdDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hdDMLP-1 production, and in vaccines or for replacement therapy. The conduction, and in vaccines or for replacement therapy. The disorder associated with the expression of hdDMLP-1, in particular heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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21-SEP-2000;
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2000GB-00024263
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RESULT 2798
ABNO1546/c
ID ABNO154
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XX Human G
XX Human;
XW muscle;
XW muscle;
XW muscle;
XW muscle;
XW muscle;
XX Mo20019
XX O6-DEC-
XX 21-SEP-
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PR 30-JAN-

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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to
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04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                               Disclosure;
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2000US-023458PP.

2000US-0236359P.

2001WO-US000661.

2001WO-US000661.

2001WO-US000663.

2001WO-US000664.

2001WO-US000665.

2001WO-US000667.

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2001WS-0266860P.
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Pred. No.
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cc protein variants having desired phenotypic improvements, and for cxxpressing the proteins. The hGDMLP-1 proteins or polypeptides may be cc used as immunogens to raise antibodies that specifically recognise hGDMLP cc -1 proteins, as standards in assays used to determine the concentration cc and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as ctherapeutic supplement in patients having specific biomolecule cc production, and in vaccines or for replacement therapy. The cc polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a cc disorder associated with the expression of hGDMLP-1, in particular heart can skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. Cc The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. Cc The sequence data for this patent did not form part of the printed cc specification, but was obtained in electronic format directly from WIPO cc at ftp.wipo.int/pub/published_pct_sequence
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ABN01545/c
ID ABN01545 standard; DNA; 17
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine;
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                                                                     AEOMICA
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                                                                                             2000US-0207456P.
2000US-023458PP.
2000US-0236359P.
2000US-0236359P.
2001WO-US000661.
2001WO-US000662.
2001WO-US000664.
2001WO-US000665.
2001WO-US000667.
2001WO-US000668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 22; gene disorder; amplicon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            scanning SEQ ID NO:4 sequence
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Pred. No.
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1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hGDMLP-1; heart;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ID NO:1537.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ç
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cc nucleic acids can be used as probes to detect, characterise and quantify cc hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 cc protein variants having desired phenotypic improvements, and for c expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 cused as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 groduction, and in vaccines or for replacement therapy. The colymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a colymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a colymucleotide sequence encoding hGDMLP-1, in particular heart col and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the present sequence in the exemplification of the present invention. N.B. of the present invention. N.B. of the present invention of the printed capture of the present invention.
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Best Local S
Matches 15
                                                                                                                                                                                                                                  (GENO-)
(NIGE-)
(EISA)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ss; PCR; acetyltransferase; 20-90-05; allergic disease; primer; differential display; eosinophil; antiallergic; atopic dermatitis; GT1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 1537; 214pp; English
                                                                         WPI; 2002-315738/35
                                                                                                                                                                                                                                                                                                                                                                 25-SEP-2000; 2000JP-00291318.
                                                                                                                                                                                                                                                                                                                                                                                                                                21-SEP-2001; 2001WO-JP008246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK49634;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK49634 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Acetyltransferase-like
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                                                                                                                                                                                                                                  RES INC.
GEN NAT CHILDREN'S
CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein 20-90-05 PCR primer GT15A
                                                                                                                                                                                                                                                                       HOSPITAL
                                                                                                                                                                      Fujishima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiallergic; atopic dermatitis; GT15A
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                                                                                                                                                                   Nagasu
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                                                                                                                                                                      Taujimoto
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Sequence

17

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U; 0 Other;

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(i.e. differential display). Also included are methods of screening for candidate compounds which affect the expression level of the gene or the activity of the protein encoded by the gene (including related proteins and mutants), the use of probes based on the gene sequence in the examination of allergic diseases, the use of reporter constructs in the screening of candidate compounds, a vector containing a the transcription controlling region of the gene, cells transformed with the vector, an antibody against the protein and a model animal for allergic diseases which is a transgenic non-human vertebrate with lowering of expression intensity of the gene in eosinophils. The method is examining allergic diseases particularly atopic dermatitis which is also applicable in screening candidate compounds for remedies. Such method can be performed in high throughput, at low cost. The present sequence is a differential display PCR primer for the cDNA encoding the human acetyltransferase-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        different expression particularly increased expression in remission stage, in eosinophils of patients, also applicable in screening candidate compounds for remedies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method for examining allergic diseases comprises determining the expression level of a gene containing, the human cDNA appearing as ABK49633 which has homology with acetyltransferases in the eosinophils of a patient and comparing the expression level with that in the eosinophils of a healthy individual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 56; 72pp; Japanese
                                                          20-90-05
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Query Match Best Local : Matches 4464 TTTTTTTTTTTTTT 4478 15; N Similarity TTTTTTTTTTTTTT 16 Conservative (0.2%; Score 15; Pred. No. <u>.</u> Mismatches DB 1; 1 Length 17; Indels 0; Gaps

0

RESULT 2801 ABK49635 Human Acetyltransferase-like protein 20-90-05 PCR ABK49635; 15-JUL-2002 ABK49635 standard; DNA; 17 (first

Human; ss; PCR; acetyltransferase; 20-90-05; allergic disease; primer; differential display; eosinophil; antiallergic; atopic dermatitis; GT15C primer GT15C

Homo sapiens.

WO200224903-A1

28-MAR-2002

21-SEP-2001; 2001WO-JP008246.

25-SEP-2000; 2000JP-00291318.

(GENO-) GENOX RES INC (NIGE-) JAPAN GEN NAT (EISA) EISAI CO LTD. CHILDREN'S HOSPITAL

Sugita Y, Takahashi Hashida R, B; Ogawa ~ Fujishima T, Nagasu Ή Tsuj imoto ູດ

WPI; 2002-315738/35.

Examining allergic diseases by differential different expression particularly increased in eosinophils of patients, also applicable display of gene showing expression in remission in screening candidate stage

Examining allergic diseases by

differential display of

gene showing

Example 1;

Page 56; 72pp;

Japanese.

compounds for remedies

#X#X55555555555555555555555555555555

Query Match Best Local

0.2%;

Score 15; DB 1; Lo Pred. No. 1.5e+03;

Length 17;

0

Local

Sequence 17 BP; 0

A;

1 C; 1 G; 15 T; 0 U; 0 Other;

PCR primer for 20-90-05

The invention relates to a method for examining allergic diseases CC comprises determining the expression level of a gene containing, the CC human cDNA appearing as ABK49633 which has homology with CC candidate compounds which at in the eosinophils of a patient and comparing the Expression level with that in the eosinophils of a healthy individual CC (i.e. differential display). Also included are methods of screening for CC candidate compounds which affect the expression level of the gene or the CC activity of the protein encoded by the gene (including related proteins CC and mutants), the use of probes based on the gene sequence in the CC examination of allergic diseases, the use of reporter constructs in the CC examination of allergic diseases, the use of reporter constructs in the CC examination of allergic diseases, the use of reporter constructs in the CC examination of allergic mon-human vertebrate with lowering of expression CC controlling region of the gene, cells transformed with the vector, and controlling region of the gene, cells transformed with the vector, and controlling region of the gene, cells transformed with the vector, and controlling region of the gene, cells transformed with lowering of expression CC diseases particularly atopic dermatitis which is also applicable in CC diseases particularly atopic dermatitis which is also applicable in the protein 20-00.05 candidate compounds for remedies. Such method can be performed control 20-00.05 candidate compounds for semestic sequence is a differential CC display PCR primer for the CDNA encoding the human acetyltransferase-like control 20-00.05 candidate compounds for semestic control 20-00.05 candidate compounds for semestic control 20-00.05 candidate compounds for semestic candidate compounds candidate compounds for semestic candidate compounds candidate compounds for semestic candi

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RESULT 2802
ABL59038
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The specification describes a method for examining allergosis. The method comprises measuring the expression level of the gene given in ABL59037, and comparing it with the expression level of the gene in the eosinophils of a healthy person. The method is used for the examination of
                                                                                                                                                                                                                                                                                                                Nucleotide sequence of PCR primer GT15A.
                                                                                                                                                                                                                                                                                                                                                                                   ABL59038 standard; DNA; 17
                                                                            Examining allergosis, specific gene, and cohealthy control.
                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                       20-AUG-2002
                                                                                                                                                                                                                                                                                                                                                             ABL59038;
                                                                                                                                                                                                                                                                                           Human;
                                                      Example 1;
                                                                                                                         WPI; 2002-439993/47.
                                                                                                                                                                               25-SEP-2000; 2000JP-00291316.
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                                                                                                                                                                                                                                               JP2002095500-A
                                                                                                                                                          (GENO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              4464 TITTTTTTTTTTT 4478
                                                                                                                                                                                                                                                                                           allergosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 Similarity
15; Conserv
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                                                                                                                                              GENOX SOYAKU KENKYUSHO KK. KOKURITSU SHONI BYOIN INCHO.
                                                      Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                    2000JP-00291316.
                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                     17;
                                                                                                                                                                                                                                                                                           eosinophil; PCR; primer;
                                                                                       comparing
                                                      20pp;
                                                                                       involves measuring the expression levels of a mparing it to the levels in the eosinophils of
                                                                                                                                                                                                                                                                                                                                                                                                                                          16
                                                      Japanese
                                                                                                                                                                                                                                                                                                                                                                                   ВP
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RESULT 2804 ABN99829

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standard;

DNA;

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15-AUG-2002 ABN99829; ABN99829

(first entry)

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RESULT 2803
ABL59039
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The specification describes a method for examining allergosis. The method comprises measuring the expression level of the gene given in ABL59037, and comparing it with the expression level of the gene in the eosinophils of a healthy person. The method is used for the examination of allergosis. The present sequence represents a PCR primer, which is used allergosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 17; 20pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Examining allergosis, involves measuring the expression levels of specific gene, and comparing it to the levels in the eosinophils of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-439993/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence of PCR primer GT15C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-AUG-2002
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1.5e+03;
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                                                                                                                            Human; allergy;
                                                                                                                                                 Human allergic disease related
                                                                                                                                                                        15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Method for examining allergic diseases by differential display of seventeen genes showing different expression particularly significant increase in essinophils in patients with mild atopic dermatitis, also
          13-OCT-2000; 2000JP-00314093
                              28-SEP-2001; 2001WO-JP008574.
                                                                         WO200233069-A1
                                                                                             Homo sapiens.
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Matches 15; Conserv
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                                            Examination of allergic diseases comprises detecting gene B1153 (expressed in T cells of allergy patients for diagnosis treatment investigation of atopic skin inflammation and asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          eosinophils in a patient and comparing the expression level with that in the eosinophils of a healthy individual. The method can be used to examine allergic diseases, particularly atopic dermatitis, and its early diagnosis, which is also applicable in screening candidate compounds for remedies. The present sequence is a PCR primer described in the
                                                                                             WPI; 2002-713252/77.
                                                                                                                                            (GENO-)
                                                                                                                                                                               21-DEC-2000; 2000JP-00389476.
                                                                                                                                                                                                      21-DEC-2001;
                                                                                                                                                                                                                                                                                                                 Human; allergy; B1153; differential expression; antiallergic; antiasthmatic; antiinflammatory; atopic skin inflammation; PC
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The present invention relates to a method

of examining allergic

diseases

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; allergy; B1153; differential expression; antiallergic; asthma; antiasthmatic; antiinflammatory; atopic skin inflammation; PCR; prime
                                                                                                                                          The present invention relates to a method of examining allergic diseases which comprises comparing the expression level of gene B1153 in allergy patients with the expression level in healthy subjects. The method is useful for the treatment, prevention, diagnosis and study of allergic diseases including atopic skin inflammation and asthma. The present sequence is a PCR primer described in the exemplification of the
                                                                                                                                                                                                                                                               Examination of allergic diseases comprises detecting gene expressed in T cells of allergy patients for diagnosis tre investigation of atopic skin inflammation and asthma.
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Pred. No.
                                                               Score 15;
Pred. No.
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RESULT 2809
AAL47235
ID AAL4723
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Best Local S
Matches 15
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Takahashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method for examining allergic diseases with intersectin 2 gene or a gene with equivalent function of intersectin 2 as an indicator gene, which comprises determining the expression level of the gene in the eosinophils in a patient, and comparing the expression level with that in the eosinophils of a healthy individual. The method is for examining allergic diseases, particularly atopic dermatitie, which is also applicable in screening candidate compounds for remedies. The present sequence is an anchor primer described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Method for examining allergic diseases by differential display intersectin 2 gene showing different expression particularly si increase in eosinophils in patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Allergic disease; atopic dermatitis;
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                                                                                                      Allergic disease
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(NIGE-) JAPAN GEN NAT
(EISA) EISAI CO LTD.
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dermatitis; human; PCR; primer; ss.
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                                                                                                      examination method related anchor primer SEQ ID NO: 3.
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Pred. No. 1.5e+03
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Allergic disease; allergy; antiallergic; atopic dermatitis; human; PCR; primer; se

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RESULT 2810
ABK57743
ID ABK5774
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Best Local
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                                                                                                                                                                                                                                                                                                                                           Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinet
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Takahashi
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(NIGE-)
(EISA)
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  (RIBO-) RIBOZYME PHARM INC
                                                                                                      09-AUG-2001;
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GEN NAT CHILDREN'S HOSPITAL.
CO LTD.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to enzymatic nucleic acid molecules that down cregulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, chance, are useful for treatment of a patient having a condition cassociated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to cates the condition and cell. This sequence represents an acetylcysteine conditions of the coll to a cell of the discased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
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                                                                                                                                                                                                                                                                   Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinet.
               (RIBO-) RIBOZYME PHARM INC. (SYNT ) SYNTEX USA LLC. (THOM/) THOMPSON J.
                                                                                  09-AUG-2000; 2000US-0224383P
                                                                                                                09-AUG-2001; 2001WO-US024970
                                                                                                                                                     14-FEB-2002.
                                                                                                                                                                                      WO200211674-A2
                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                     acetylcysteine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK57744 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic polynucleotide that down regulates expression of channel calcium activated gene, useful for treating Chronic pulmonary disease (COPD), chronic bronchitis and asthma.
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                                                                                                                                                                                                                                                                                                                                                       gene enzymatic nucleic acid #2115.
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Pred. No. 1.5e+03;
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onic obstructive
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RESULT 2812
ABK49757
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Matches 13
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         Examining allergic diseases by differential display of genes showing different expression particularly increase in remission stage in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                     WPI; 2002-330097/36.
                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                     Atopic dermatitis; allergic disease; a
                                                                                                                                                                                                                                                                                                                           Human atopic
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JAPAN
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                                                                                                                                                   2000JP-00293021.
                                                                                                                                                                                                                                                                                                                             dermatitis cDNA related PCR primer GT15c.
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                                                                                                            CHILDREN'S HOSPITAL.
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Pred. No. 1.
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                                                                                Fujishima
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RESULT 2813
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Best Local
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            This invention relates to gene sequences that are differentially expressed in cosinophils from patients with atopic dermatitis in the expressed in ecsinophils from patients with atopic dermatitis in the increment stage as compared with those in the remission erage. These sequences are used in a novel method for examining allergic diseases comprising determining the expression levels of these genes and comparing the expression level with that in the eosinophils of a healthy individual. The method of the invention may have antiallergic or dermatological activities. The method can be used to disgnose allergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           candidate compounds for remedies. The method of the invention can be performed in high throughput, at low cost. The present sequence represents the GTISC PCR primer used to amplify the differentially amplified atopic dermatitis related cDNA sequences of the invention
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                                                                                                                                                                        Example 1; Page 54; 74pp; Japanese
                                                                                                                                                                                                           Examining allergic diseases by differential display of genes showing different expression particularly increase in remission stage in eosinophils in patients.
                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                  Sugita Y,
                                                                                                                                                                                                                                                                                                                                                                                                        26-SEP-2000; 2000JP-00293021
                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-2001; 2001WO-JP008247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human atopic dermatitis cDNA related PCR primer GT15a.
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                                                                                                                                                                                                                                                                                                                   Hashida R,
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antiallergic; dermatological; GT15e
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                                                                                                                                                                            This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease
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                                                                                                                                                                                                                                                                Claim 1; Page
                                                                                                                                                                                                                                                                                      New nucleic acid sequences associated with tumor suppression, regres apoptosis or virus resistance are useful to diagnose and treat viral disease, development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human tumour suppressor sequence #1412.
                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                               20-JUN-2001; 2001FR-00008139.
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                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 BP; 1 A; 0 C; 1 G;
                                                                       ATCCCAGCTGGGACC 2409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTTTTTTTTTTTTT 16
                                                                                                                                                                                                                                                                                                                                                                Telerman A,
                                                                                              Conservative
                                                                                                                                             BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                               366; 798pp; French.
                                                                                                                                                                  development
                                                                                                                                             ₽
                                                                                                                                                                                                                                                                                                                                                                                        ENGINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                apoptosis; virus
                                                                                                        100.0%;
                                                                                                                                           6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antitumour; cytostatic; tumour suppression;
ptosis; virus resistance; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
                                               16
                                                                                                                                                                                                                                                                                                                                                                                         LAB
                                                                                                                                                                                                                                                                                                                                                                 Amson R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                              0
                                                                                                         Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No.
                                                                                                                                                                   of tumour
                                                                                                                                                                                                                                                                                                                                                                                         SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15
                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T;
                                                                                                                                                                   cells or cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     U; 0 Other;
                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                         1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.5e+03;
                                                                                                                   Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                  degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                regression,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                              Gaps
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RESULT 2816
ABT36431/c
ID ABT3643
XX
AC ABT3643
XX
DT 12-JUNXX

ABT36431

standard;

DNA;

17

BP

12-JUN-2003 ABT36431;

(first entry)

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Matches
                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                       comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenaemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX79793;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST; expressed sequence tag; ss; polymorphic repeat; tandem rej polymorphic marker prediction of ubiquitous simple sequences; lepy, human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying a candidate polymorphic repeat within a coding sequence, i understanding or treating genetic disease, comprises detecting tandem repeats in a target coding sequence and scoring the repeats for polymorphic probability.
                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Garner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EST polymorphic DNA repeat polynucleotide #118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         repeat within a coding sequence (expressed sequence tag, EST),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention discloses a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6472154-B1
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                                                  4464 TTTTTTTTTTTTT 4478
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                                                                                                     15;
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۳
                                                                                                                              Similarity
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                                                                                                                                                                                                             BP; 0 A;
                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       483; 588pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-00475947.
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                                                                                                                           0.2%;
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                                                                                                                                                                                                             2 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Minna JD,
15
                                                                                                  0,
                                                                                                                              Score 15;
Pred. No.
                                                                                                                                                                                                          15 T; 0
                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for identifying a candidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fondon JW
                                                                                                                                                                                                          U; 0 Other;
                                                                                                                              1.5e+03;
                                                                                                                                                       DB 1;
                                                                                                                                                       Length 17;
                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tandem repeat;
                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POMPOUS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           which
                                                                                                     Gaps
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Tumour suppression related human fukutin oligo SEQ ID

No 2068

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ADB04274
ID ADB0
XX ADB0
AC ADB0
XX DT 20-
XX
DE Hum
XX
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                                                                                                                                                                                                                                                                                       8
                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel isolated 17 mer nucleic acid sequence, cc given in the specification, a sequence containing at least 15 consecutive cc alignment, at least 80 % identity to the 17 mer sequence with, after optimal cc alignment, at least 80 % identity to the 17 mer sequence, a sequence that cc hybridizes to them under highly stringent conditions, or the complement cc fany of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, cc identifying, quantifying and/or amplifying a nucleic acid, e.g. as one cc component of a gene chip, in vitro as (anti)sense reagents, and for cromponent of a gene chip, in vitro as (anti)sense reagents, and for component of a gene chip, in vitro as (anti)sense reagents, and for cromponent of a gene chip, in vitro as (anti)sense reagents, and for component of a gene chip, in vitro as (anti)sense reagents, and for cromponent of a gene chip, in vitro as (anti)sense reagents, and for component of a gene useful for component of pharmaceuticals for prevention and/or treatment of viral cd diseases that are characterised by development of tumours or cell containing the action and pharmaceuticals for prevention and/or treatment of viral cd degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in cation sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene chips.
                                                                                                                                                                                        RESULT 2817
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 274; 720pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   with tumors and cell d
and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-313353/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-2001; 2001FR-00011978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-2002; 2002WO-IB004208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                       Human MDZ7 scanning oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 3
                                                                 20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                               ADB04274 standard; DNA; 17
                                                                                                                                                                                                                                                                        6294 CTGGCCTCCAGGGAT 6308
                                                                                                                                                                                                                                                   16
                                                                                                                                                                                                                                                                                                                                 l Similarity
15; Conserv
                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amson R,
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tuijnder M;
                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                  Score 15; DB 1; L
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                   Mismatches
                           SEQ
                           Ħ
                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                       Length 17;
                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                   Gaps
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PCR; primer;

88.

01-JAN-2004

(first entry

ADC84469;

ADC84469 standard;

DNA; 17

Plant blastogenesis; transformation; gene expression; tissue specific; PCR primer for amplifying plant blastogenesis specific gene #SEQ

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03-JUN-2003

JP2003159071-A

밁 8

Matches Query Match Best Local

l Similarity
15; Conservat

Conservative

0

100.0%; 0.2%;

Score 15; DB 1; Pred. No. 1.5e+0 Mismatches

1.5e+03;

0,

Gaps

0

Length 17; Indels

4470 TITTTTTTTTTTTG 4484

RESULT 2818

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WANTE CONTROL OF THE 
                                                                                                                                                                                                                              proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 1p21.3-22., MDZ7 is encoded at chromosome 1p21.3-22., MDZ7 is encoded at chromosome 1p21.3-2 and MDZ12 is encoded at chromosome 1p21.2 and MDZ12 is encoded at chromosome 1p21.2 and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ1, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 gross are alterations in MDZ3, MDZ4, MDZ7, or MDZ12 acids can also be used as probes to detect and characterize gross are alterations in MDZ3, MDZ4, MDZ7, or MDZ12 gross are more detect.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21:3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUL-2002; 2002EP-00016874
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Sequence 17 BP; 1 A; 0 C; 2 G; 14 T; 0 U; 0 Other;
                                                                                       useful in constructing microarrays for measuring gene expression. proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the inventio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; SEQ ID NO 5260; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-423107/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-AUG-2001; 2001US-00922181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ନ୍ଧ Y,
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RESULT 2819
ADC8446
ADC8446
AC ADC846
AC ADC8466
AC ADC8466
AC ADC8466
AC ADC8468
AC ADC8668
AC ADC
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
  The invention relates to naturally derived DNA specifically expressed during plant blastogenesis. The DNA of the invention is useful for producing a transformed plant. Methods of the invention are also useful for compulsive expression of this DNA. Methods of the invention are useful for plant tissue specific expression of genes. Also, the growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to naturally derived DNA specifically expressed during plant blastogenesis. The DNA of the invention is useful for producing a transformed plant. Methods of the invention are also useful for compulsive expression of this DNA. Methods of the invention are useful for plant tissue specific expression of genes. Also, the growth stage of a plant can be controlled specifically. The current sequence represents a PCR primer for amplifying a plant blastogenesis specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New naturally derived DNA specifically expressed during blastogenesis of a plant, useful for producing a transformed plant and for compulsive expression of a protein.
                                                                                                                                                                                                                                                      New naturally derived DNA specifically expressed during blastogenesis a plant, useful for producing a transformed plant and for compulsive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP2003159071-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC84468 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3;
                                                                                                                                                                                                                                                                                                                                          WPI; 2003-818678/77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Plant blastogenesis; transformation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for amplifying
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                                                                                                                                                                            Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-NOV-2001; 2001JP-00358366.
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                                                                                                                                                                                                                                                                                                                                                                                                DOKURITSU GYOSEI HOJIN
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                                                                                                                                                                      SEQ ID NO 1; 43pp; Japanese.
                                                                                                                                                                                                                                of a
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                                                                                                                                                                                                                             protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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ADE77745
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         stage of a plant can be controlled specifically. The current represents a PCR primer for amplifying a plant blastogenesis gene of the invention.
                                             This invention relates to a novel method for screening and examining allergic diseases by the use of B1799 as the indicator gene. Specifically, it comprises determining the expression level of this indicator gene in a biological sample obtained from the patient, and identifying differential expression (increased expression of B1799) in comparison to that observed in a healthy individual. The present invention describes the B1799 protein as antiallergic, antiinflammatory and dermatological. As such, through the use of gene therapy, this method can be used to treat allergic diseases particularly atopic dermatitis. Furthermore, it is useful for determining a diagnosis that is convenient and non-invasive, and is also applicable in high throughput screening to identify candidate compounds for additional remedies. This
                                                                                                                                                                                                                                           Examining allergic diseases, such as atopic dermatitis, comprises comparing the expression levels of gene B1799 in T cells in a pata healthy individual.
                                   identify candidate compounds for addoligonucleotide sequence is the DNA
                                                                                                                                                                                                                  Example 1; SEQ ID NO 5; 87pp; Japanese.
                                                                                                                                                                                                                                                                                              WPI; 2003-804076/75.
                                                                                                                                                                                                                                                                                                                       Matsumoto
                                                                                                                                                                                                                                                                                                                                                                                     03-APR-2002; 2002JP-00100908
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; allergic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                 (NIGE-)
                                                                                                                                                                                                                                                                                                                                                             (GENO-)
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                                                                                                                                                                                                                                                                                                                       Yoshida
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Pred. No.
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RESULT 2822
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Matches 15; Conserv
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Best Local
                                                                                                                                                                                                                                                                                    This is the nucleotide sequence of a PCR primer used in the me invention, involving the use of novel apoptosis-related DNAs a proteins. The inventions can be used as diagnostic reagents fo e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can a for treatment of apoptosis-related diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV54175 standard;
                                                                                                                                                                                                                                                                    Sequence 18 BP; '0 A;
                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 51; 70pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                            treating
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                                                PCR; primer; amplifimmunohistological
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            WO9839437-A1
                               Synthetic
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                                                                                                                      AAV54173
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                                                                                                                                                                                                                                                                                                                                                                            apoptosis-related DNAs ing diseases associated
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Pred. No.
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                                                           antibody; inhibition, 88;
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DNAs and
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel apoptosis-related DNAs and proteins - for diagnosis, preventing treating diseases associated with apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sakaki Y;
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                                                                                                                                                                                                                                                             Nucleotide sequence PCR primer
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                                                                                      WPI; 1998-495844/42
                                                                                                         Sakaki
                                                                                                                                            05-MAR-1997;
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                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                   PCR; primer; amplification; immunohistological staining.
                                                                                                                                                                                                                                                                                21-DEC-1998
                                                                                                                                                                                                                                                                                                 AAV54164;
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                                                                                                                                                                                                                                            apoptosis; antibody; inhibition; ss;
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Pred. No.
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1.6e+03;
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This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in

Novel apoptosis-related DNAs and proteins - treating diseases associated with apoptosis.

HOT

diagnosis,

preventing

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Example

1;

Page 47;

70pp;

Japanese

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RESULT 2824
AAV54166
ID AAV5416
XX AAV5416
XC AAV5416
XC AAV5416
XX PCR; pr
KW PCR; pr
KW immunoh
XX Synthet
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RESULT 2825
AAZ90649
ID AAZ9064
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AC AAZ9064
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AC AAZ9064
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Matches 15
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Matches 15; Conserv
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13-JUN-2000
                                                                                                                                                               This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also be used for treatment of apoptosis-related diseases
                  AAZ90649;
                                                                                                                                                 Sequence 18
                                                                                                                                                                                                                                             Novel apoptosis-related DNAs and proteins - treating diseases associated with apoptosis.
                                                                                                                                                                                                                                                                        WPI; 1998-495844/42.
                                                                                                                                                                                                                                                                                                                                                              11-SEP-1998
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                                 AAZ90649
                                                                                                                                                                                                                            Example 1; Page 48; 70pp; Japanese.
                                                                                                                                                                                                                                                                                                                           05-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                           (KYOW ) KYOWA HAKKO KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                              WO9839437-A1
                                                                                                                                                                                                                                                                                                                                                                                                                PCR; primer; amplification; apoptosis; antibody; inhibition; ss; immunohistological staining.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleotide sequence PCR primer 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunohistological staining, as apoptosis inhibitors. It can also be used for treatment of apoptosis-related diseases
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                                                                                                              l Similarity
15; Conserv
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                                 standard;
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Pred. No.
                                                                                                                                                15 T; 0 U; 0 Other;
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hes 0;
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RESULT 2826
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to identification of genes and proteins of adipose tissue relating to obesity, particularly complications of visceral obesity including diabetes, hypertipemia, hypertension, arteriosclerosis, hyperticemia and sleep apnea syndrome. The genes (AAZ30631-633) and the proteins (AAY67598-Y67600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ30640-51
                                                                                                                            23-JUL-1998;
                                                                                                                                                                         08-FEB-2000.
                                                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                        arteriosclerosis;
                                                                                                                                                                                                                                                                                                                        Adipose tissue; obesity; diabetes; hyperlipemia; hypertension; human;
                                                                                                                                                                                                                                                                                                                                                                   Human adipose tissue gene amplifying primer #9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ90648;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 1 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A physiologically active protein specifically derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                 23-JUL-1998;
                                                                                                                                                                                                                  JP2000037190-A
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                                          (NISB ) JAPAN TOBACCO
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2000-306578/27.
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15; Conserv
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                                                                                 98JP-00225228
                                                                                                                              98JP-00225228
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                                                                                                                                                                                                                                                                                                      hyperuricemia; sleep apnea syndrome; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amplifying the human adipose
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Pred. No.
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Example 2; Page 18; 50pp; Japanese

A physiologically active

protein specifically derived from mammal tissue

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RESULT 2828
AAZ90651
ID AAZ9065
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AAZ90646
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 1 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                               tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hypertransic comits and sleep apnea syndrome. The genes (AAZ90631-633) and the proteins (AAX67598-457600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ90640-51 represent PCR primers amplifying the human adipose tissue genes
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-306578/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Adipose tissue; obesity; diabetes; hyperlipemia; hypertension; human; arteriosclerosis; hyperuricemia; sleep apnea syndrome; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human adipose tissue gene amplifying primer #7.
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                                                                                                                                                                                       Sequence 18 BP; 2
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                                                                                                                                                                                                                                                                                                    The invention relates to identification of genes and proteins of adipose
                                                                                                                                                                                                                                                                                                                                                                                                                     (NISB ) JAPAN TOBACCO
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                                                                                                                                                                                       A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                            active protein specifically derived from mammal tissue
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100.0%; Pred. No.
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                                                                                                                                             Score 15;
Pred. No.
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1.6e+03;
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AAZ90651 standard; DNA; 18 BP

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AAA58385/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A physiologically active protein specifically derived from mammal tissue.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    arteriosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 0 A;
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arteriosclerosis; hyperuricemia; sleep apnea syndrome; PCR primer; ss.
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Barbera-Guillem E,
                                             10-NOV-1998;
                                                                                                                                                       Synthetic
                                                                                                                                                                              Nanocrystal;
                                                                                                                                                                                                    Polynucleotide # 1 used in a biomolecule detection system.
                                                                                                                                                                                                                             01-NOV-2000
                                                                                                                                                                                                                                                                        AAA58385
                                                                                 10-NOV-1999;
                                                                                                         18-MAY-2000
                                                                                                                                WO200028088-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       represent PCR primers amplifying the human adipose tissue
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                      (BIOC-)
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                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                              biomolecule detection; nonisotopic detection system; ss.
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                                             98US-0107828P.
99US-00437076.
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                                                                                 99WO-US026612
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                                                                                                                                                                                                                                                                          DNA; 18
 Nelson MB,
                                                                                                                                                                                                                                                                                                                                                                                                                                2 C; 1 G; 15 T; 0 U; 0 Other;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to functionalised nanocrystals for use in nonisotopic detection systems for biomolecules e.g. nucleic acids, proteins, lipids or drugs. The nanocrystals have polynucleotide strands attached to their surfaces with one end of the polynucleotide extending outwardly from the nanocrystal. The present sequence is one such polynucleotide. These nanocrystals are used with a second series of nanocrystals, which have polynucleotides complementary to the first polynucleotides, so that the respective complementary strands hybridise to each other and form a dendrimer. This dendrimer produces a signal which can then be detected e.g. fluorescence. The present sequence is composed mainly of Adenine bases. This sequence may therefore be used with a polynucleotide composed mainly of Thymine bases (AAAS8386)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Functionalized nanocrystal carrying polynucleotide, target analyte, forms dendrimers with complementary amplify the fluorescent signal.
                                                                                                                      Functionalized nanocrystal carrying
                                                                                                                                                                      Barbera-Guillem
                                                                                                                                                                                                                     10-NOV-1998;
09-NOV-1999;
                                                                                                                                                                                                                                                                              18-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                          Polynucleotide # 2 used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA58386 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                        10-NOV-1999;
                                                                                                                                                                                                                                                                                                      WO200028088-A1
                                                                                                                                                                                                                                                                                                                                                    Nanocrystal;
                                                                                                                                                                                                                                                                                                                                                                                                                           AAA58386;
                                                                                                                                                                                              (BIOC-) BIOCRYSTAL
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/ the fluorescent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                   biomolecule detection; nonisotopic
                                                                                                                                                                                                                                                                                                                                                                                                  (first
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99US-00437076.
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                                                                                                          nanocrystal carrying polynucleotide, forms dendrimers with complementary
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                                                                                                                                                                                                                                                                                                                                                   detection system;
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                                                                                                            nanocrystals
                                                                                                                      used for detecting
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The present invention relates to functionalised nanocrystals for nonisotopic detection systems for biomolecules e.g. nucleic acid proteins, lipids or drugs. The nanocrystals have polynucleotide attached to their surfaces with one end of the polynucleotide exoutwardly from the nanocrystal. The present sequence is one such

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extending strands

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Example 3;

Page

69; 72pp;

English.

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RESULT 2831
AAF26668
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Best Local S
Matches 15
        The present invention describes an antisense compound (I) of up to 30 nucleobases in length capable of inhibiting the expression of human Smad7. (I) has antiinflammatory and cytostatic, and is a modulator of Smad7 expression. (I) can be useful for inhibiting the expression of human Smad7 in human cells or tissues, in vitro. (I) is commonly used a research research and in diagnostics for example, to elucidate the function of particular genes. (I) is also useful for distinguishing between functions of various members of a biological pathway and for research use. (I) is also useful for diagnostics, therapeutics, prophylaxis and in kits. (I) is also useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. AAF26667 AAF26706 represent human Smad7 antisense oligonucleotides from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polynucleotide. These nanocrystals are used with a second series of nanocrystals, which have polynucleotides complementary to the first polynucleotides, so that the respective complementary strands hybridise to each other and form a dendrimer. This dendrimer produces a signal which can then be detected e.g. fluorescence. The present sequence is composed mainly of Thymine bases. This sequence may therefore be used with a polynucleotide composed mainly of Adenine bases (AAA58385)
                                                                                                                                                                                                                           Antisense compound cap
useful for preventing
                                                                                                                                                                                                                                                                       WPI;
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Elammatory; cytostatic; infection;
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inflammation; tumour
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our formation;
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RESULT 2832
AAS17036/c
ID AAS17036
XX AAS1703
XX AC AAS1703
XX Human G
XX Lucifer
XX Lucifer
XX Examcer;
XX Cancer;
XX MO20018
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The invention relates to a yeast cell comprising nucleotide sequences cc encoding first and second heterologous fusion proteins comprising first cand second peptides (P1, P2) of a peptide binding pair joined to a ctranscriptional activation protein (TAP) DNA binding domain or TAP cc transcriptional activation domain, respectively, and a luciferase gene cc activated by positive transcriptional control of TAP reconstituted by cobinding of P1 and P2. The cell is useful for detecting the interaction of a first peptide and a second peptide of a peptide binding pair which cc involves culturing the cell, incubating a test sample with the yeast cell cunder conditions suitable to detect the selected phenotype and detecting the level of expression of the luciferase gene. The cell is also useful cc for determining whether a test sample interacts with a first or second peptide of a peptide binding pair. The yeast cell comprises a nucleotide concernment of the peptide strategiant mimics and receptor mimics. The cell may be used for monitoring the binding of peptides by peptide binding pair compounds interactions, e.g., hormone/receptor conteractions. The cells are useful in the detection of the ability of the test sample to affect the binding of a peptide binding pair and contend through cells are also useful in the detection of the ability contends interaction. The screening methods can be used confidentifying compounds interacting with any peptide binding pair and contends interaction of that interaction, e.g., protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New yeast cell, useful expresses heterologous binding pair joined to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-049273/06
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transcriptional
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Page 53; 119pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for determining protein-protein interactions, fusion proteins comprising peptides of peptide transcriptional activation protein DNA binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             o
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RESULT 2833
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ID AAA83021 standard; DNA; 19
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                                        Matches
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                                                                                                                             The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinas other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient
                                                                                                                                                                                                                                                          New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                         Tritz
                                                                                                                                                                                                                                                                                                                                                                                                                    06-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA83021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18
                                                                                        Sequence 19
                                                                                                                   restenosis treatment
                                                                                                                                                                                                                                   Disclosure; Page 55; 109pp; English.
                                                                                                                                                                                                                                                                                                                 WPI; 2000-412314/35
                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cdk6 ribozyme binding site #81.
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                                        l Similarity
15; Conserv
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TTCTGCAAATATGCA 1692
                                                                                                                                                                                                                                                                                                                                         Welch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hairpin;
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                                                                                                                                                                                                                                                                            for inhibiting restenosis, cleaves dependent kinase other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy; vasotropic; restenosis;
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                                                                 DB 1;
                                                    1.8e+03;
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                                                                                           Other;
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                                                                 Length 19;
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                                         Gaps
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                                                                                                                        nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn are account of the contractive used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                 ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                          skin
                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 116; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-OCT-2000; 2000WO-US029500.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robbins JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sickle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                        present invention describes a method for treating a proliferative or eye disease and scarring. The method involves administering a
                                                                                                              AAH57577 to AAH62099 represent sequences used lification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
 l Similarity
15; Conserv
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   Conservative
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                 0.2%; Score 15; DB 1; L
00.0%; Pred. No. 1.8e+03;
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AAT73291
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AAQ13458/c
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        primer; PCR; polymerase chain reaction; sequencing; walking;
complementary extension reaction; low redundancy; universal
                                                                                                                                                                                                                                                                                              This codon is a potentially oncogenic site and a group of mutant probes were synthesised based on single point mutations at codon 201. After PCI amplification of nucleic acid samples using specific primer pairs, these probes can be used to detect Gi(alpha)1 mutations associated with oncogenesis. See AAQ13431-Q13542. (Updated on 25-MAR-2003 to correct PI
                                          Primer 1 for
                                                                12-DEC-1997
                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                             This probe corresponds to a mutant sequence around codon 201 This codon is a potentially oncogenic site and a group of mut
                                                                                                                                                                                                                                                                                                                                                                                Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                    Method for detection of point mutation(s) in nucleic acid segments - where segments encode GTP binding protein or sub-unit and method involves amplification followed by sequence-specific probe hybridisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-FEB-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       adenylate cyclase activity inhibition; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Probe to mutant codon 201 (AAA) of G-protein Gi(alpha)1 subunit.
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05-NOV-1991
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                                                                                        AAT73291;
                                                                                                             AAT73291 standard;
                                                                                                                                                                                                                                                                                      oncogenesis.
field.)
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                                                                                                                                                                                                                              Similarity
                                                                                                                                                                      GITTTCACTTTAGTT
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                                                                                                                                                                                                                                                                                                                                                                                Page
                                                                                                                                                                                                                                                              BP; 10 A; 3 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first en
                                          pUC19 DNA amplification.
                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORP
                                                                                                                                                                                                                                                                                                                                                                               65; 69pp; English.
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                                                                                                              DNA;
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         primer;
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AAT73292
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Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   using a DNA polymerase, comprises a combination of fragment walking and DNA sequencing. DNA fragments are formed by digestion of DNA with a restriction enzyme and the targeted DNA sequence can be determined directly from the digested DNA fragments. By exploring the overlapping sequence of the determined base sequence, the overall base sequence of lengthy DNA can be determined with low redundancy without cloning or subcloning. In addition, the method can be done with commercially available universal primers or with fewer primers than required in existing methods. AAT73291-92 are primers used in determination of the DUC19 sequence. Primer extension was carried out using 16 primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Efficient sequencing of long DNA by fragment walking - with simultaneous sequencing of restriction enzyme fragment and adjacent region of intact DNA, avoids the need for cloning and requires fewer primers.
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30-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-1996;
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                                   18-SEP-1995;
30-JAN-1996;
                                                                                                                                                                                                                                                     AAT73292;
                                                                                                                                                                                                                                                                             AAT73292 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A method for DNA analysis based on a complementary extension reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 11; 50pp; English.
                                                                       17-SEP-1996;
                                                                                              09-APR-1997
                                                                                                                     EP767240-A2
                                                                                                                                             Synthetic.
                                                                                                                                                                               primer;
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           (HITA ) HITACHI LTD
                                                                                                                                                                                                                                                                                                                                                                4470
                                                                                                                                                                                                                                                                                                                                                                                      15;
                                                                                                                                                                                                                                                                                                                                        _
                                                                                                                                                                               PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                       for
                                                                                                                                                                                                                                                                                                                                                                TTTTTTTTTTTTG 4484
                                                                                                                                                                                                                                                                                                                                                                                                                                     20 BP; 1
                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                    polymerase
y extension
                                                                                                                                                                                                       puc19
                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95JP-00238141.
96JP-00013634.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96EP-00114907.
                                   95JP-00238141.
96JP-00013634.
                                                                       96EP-00114907.
                                                                                                                                                                                                                                                                                                                                                                                                                                     A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>ج</u>
                                                                                                                                                                                                       DNA amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                     2 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                              0.2%;
                                                                                                                                                                 chain reaction; sequencing; walking; reaction; low redundancy; universal
                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                     Score 15; DB; Pred. No. 1.9
                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 14 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                     U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                  1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                             Length 20
                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                    primer;
                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ð
                                                                                                                                                                                                                                                                                                                                                                                        0
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restriction enzyme and the targeted DNA sequence can be determined directly from the digested DNA fragments. By exploring the overlapping sequence of the determined base sequence, the overall base sequence of lengthy DNA can be determined with low redundancy without cloning or subcloning. In addition, the method can be done with commercially available universal primers or with fewer primers than required in existing methods. AAT73291-92 are primers used in determination of the pUC19 sequence. Primer extension was carried out using 16 primers
                                                                                                                                                                                                                                                                                                                                                                    Efficient sequencing of long DNA by fragment walking - with simultaneous sequencing of restriction enzyme fragment and adjacent region of intact DNA, avoids the need for cloning and requires fewer primers.
                                                                                                                                                                                                                                   using a DNA polymerase, DNA sequencing. DNA fra
                                                                                                                                                                                                                                                                              A method
                                                                                                                                                                                                                                                                                                                           Example 1; Page 11; 50pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-205424/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kambara H,
                                                                                                                                                                                                                                                                                   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Okano
                                                                                                                                                                                                                                      DNA analysis based on a complementary extension reaction polymerase, comprises a combination of tragment walking and ng. DNA fragments are formed by digestion of DNA with a
                                                                                                                                                                                                                                                                                                                           English.
G; 15 T; 0 U; 0 Other;
                                                                                                                                                                             9
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RESULT 2838
AAA72163
                                                                                                                             Matches
                                                                                                                                      Query Match
Best Local :
15-SEP-2003
24-NOV-2000
                           AAA72163;
                                              AAA72163
                                                                                                            4470 TITTTTTTTTTTTG 4484
                                                                                                                              15;
                                                                                                                                      Similarity
                                              standard;
                                                                                                                              Conservative
(revised)
(first entry)
                                               DNA;
                                                                                                                                     100.0%;
                                               20
                                                                                           15
                                                                                                                              0
                                                                                                                                       Score 15;
Pred. No.
                                                                                                                              Mismatches
                                                                                                                                        1.9e+03;
                                                                                                                               Indels
                                                                                                                              0
                                                                                                                              Gaps
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밁 S

Sequence 20

BP; 1 A;

1 C; 3

0.2%;

DB 1;

Length 20;

0

humanised antibody; complementarity determining region; CDR; human Pas; Fas ligand; apoptosis modulator; programmed cell death; autoimmune disease; allergy; attopy; arteriosclerosis; myocarditis; cardiomyopathy; glomerulonephritis; aplastic anaemia; panmyelophthisis; hepatitis; AIDS; graft rejection; heavy chain; sequencing primer; ss. Anti-Fas antibody; monoclonal antibody HFE7A; sapiens. FERM-BP-5828; murine; ; human Fas;

Humanised anti-Fas antibody heavy chain primer,

SEQ ID NO:93.

Mus musculus. Homo

JP2000169393-A Chimeric.

20-JUN-2000

30-SEP-1999; 99JP-00278301.

30-SEP-1998; 98JP-00276883

(SANY) SANKYO CO

2000-485645/43

Preventive or treating agent for the diseases caused by an abnormality the Fas/Fas ligand system e.g. autoimmune diseases, contains anti-Fas 'n

Example 15; Page 49; 139pp; Japanese

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AAZ38528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to compositions for the prevention or treatment or diseases caused by an abnormality in the Fas/Fas ligand system containing an anti-Fas antibody as the active component. The anti-Fas antibody is either the murine anti-human Fas monoclonal antibody HFE7A, or a humanised version of HFE7A containing identical CDRs (complementarity determining regions) to antibody HFE7A. Via its interaction with Fas, the antibody of the invention acts as a modulator of apoptosis. The compositions of the invention may therefore be used in the treatment or prevention of conditions such as autoimmune diseases, allergy, atopy, arteriosclerosis, myocarditis, cardiomyopathy, glomerulonephritis, applastic anaemia (panmyelophthisis), hepatitis, AlDS and organ graft
                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                               Microtubule associated protein 4; MAP4; real-time quantitative PCR; expression; microtubule; assembly; function; cytoskeleton; structural; dynamic; stabilisation; lattice; overexpression; p53; oncogene; cancer;
                                      09-APR-1999;
                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human microtubule-associated protein 4 (MAP4) antisense oligo #63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rejection. The present sequence represents a humanised HFE7A-derived an -Fas antibody heavy chain sequencing primer used in an exemplification the invention. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                     US5998148-A.
                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ38528;
            (ISIS-) ISIS
                                                                  09-APR-1999;
                                                                                             07-DEC-1999
                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                         hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                      chemotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ38528 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7181 GGTGGGCATGTGTGA 7195
                                                                                                                                                                                                                                                                                                                                                                               sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGTGGGCATGTGTGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative (
            PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                        tumour; drug sensitivity; antisense; therapy;
; inhibition; research; diagnostic; ss.
                                       9905-00289368
                                                                  99US-00289368.
                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                        /mod_ba
16. .20
/*tag=
                                                                                                                                                                                                                                                                                                 note=
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                                                                                                                                                note=
                                                                                                                                                                                                                 *tag=
                                                                                                                                                                                                                                                                                                                            *tag=
                                                                                                                                                                                                                                            note=
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                                                                                                                                                                                                     _base=
                                                                                                                                                                                                                                                                                                                base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                              2,
                                                                                                                                                                                                                                                                                                 "Phosphorothioate linkages'
                                                                                                                                                2,
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                                                                                                                                                                                                                                                                         σ
                                                                                                                                                methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                              OTHER
                                                                                                                                                                                                      m5c
                                                                                                                                                                                                                                            methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                          OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15; DB 1;
; Pred. No. 1.9e+(
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 4 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           on of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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CC against the gene encoding human microtubule-associated protein 4 (MAP4).

CI Inhibition of MAP4 expression was measured by determination of MAP4 mRNA

CC levels in a variety of cell lines via real-time quantitative PCR. The

CC cell lines used included the bladder carcinoma cell line T-24, the human

CC lung carcinoma cell line A549, human neonatal dermal fibroblasts and

CC croup of proteins that mediate microtubule-associated proteins comprise a

CC group of proteins that mediate microtubule assembly and function which is

CC required for cytoskeletal integrity. MAP4 is a member of the non-neuronal

CC structural MAP family and is believed to affect microtubule dynamics by

CC stabilising the microtubule lattice. MAP4 expression has been shown to be

CC linked to cancer chemotherapeutic drug sensitivity. These antisense

CC nolecules are useful for treating animals, particularly humans, having or

CC being prone to a disease or condition associated with the expression of

CMAP4. The oligonucleotides are also useful for research and diagnostic

CC applications
Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bennett CF, Ackermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotides for inhibiting microtubule-associated protein 4 expression, useful in treating disorders associated with microtubule protein expression.
                                                                          Sequence 20 BP; 7 A; 1 C; 11 G; 1 T; 0 U; 0 Other;
                                                                                                                             applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a preferred antisense oligonucleotide targetted against the gene encoding human microtubule-associated protein 4 (MAP4).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-052543/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Col 39; 39pp; English
0.2%; Score 15; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           된
                        Length 20
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RESULT 2840
AAA11601
                                                                                                                                             Matches
                                                                                                      AAA11601;
                                                                                                             AAA11601 standard; DNA;
                                                                                                                                     3222 TGGGAGGAGGGAAGG 3236
                                                                                                                               4
                                                                                                                                             15;
                                                                                                                                                Similarity
                                                                                                                              TGGGAGGAGGGAAGG
                                                                                                                                             Conservative
                                                                                                             20
                                                                                                                               18
                                                                                                             ВP
                                                                                                                                            ç
                                                                                                                                             Mismatches
                                                                                                                                           DB 1,
1.9e+03;
0;
                                                                                                                                             Indels
                                                                                                                                            0,
                                                                                                                                            Gaps
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Humanised HFE7A designed heavy chain DNA primer #4.

08-AUG-2000

(first entry)

닭 S

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Fas; antibody; human; anti-inflammatory; anti-anemic; antidiabetic; anti-allergic; anti-arthritic; antiviral; immunomodulatory; cardiant; dermatological; immunosuppressive; thyronimetic; antiarteriosclerotic; nephrotropic; antiinfertility; neuroprotective; antiarteriosclerotic; hepatotropic; humanized; apoptosis; systemic lupus erythematosus; Hashimoto disease; rheumatoid arthritis; graft versus host disease; Sjorgen's syndrome; anemia; Addison's disease; scleroderma; sterility; Goodpasture syndrome; Crohn's disease; sterility; myasthenia gravis; multiple sclerosis; Basedow's disease; thrombopenia purpura; allergy; insulin dependent diabetes mellitus; arteriosclerosis; myocarditis; cardiomyopathy; glomerulonephritis; hepatitis; transplant rejection;
    30-SEP-1998;
30-SEP-1998;
                                                                                                                                                                                                                                                                                 EP990663-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                             cardiomyopathy;
primer; ss.
                                                                                                                     29-SEP-1999;
                                                                                                                                                                                                     05-APR-2000
                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
98JP-00276881
98JP-00276882
                                                                                                                     99EP-00307711.
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RESULT 2841
ABL48723
  CC molecule (I) that, induces apoptosis in cells with an abnormal Fas' Fas' CC ligand system, by binding to Fas on the cell surface, and prevents CC apoptosis in cells with a normal system, by inhibiting binding between CC Fas and its ligand. The products of the invention have anti-inflammatory, CC anti-anemic, antidiabetic, anti-allergic, anti-arthritic, antiviral, CC immunomodulatory, dermatological, immunosuppressive, thyromimetic, CC antiarteriosclerotic, cardiant and hepatropic activity. (I) induce constrainteriosclerotic, cardiant and hepatropic activity. (I) induce conspicuous by binding to cell surface Fas or inhibit it by competitive continhibition of ligand binding. (I) are used to treat and/or prevent CC diseases associated with the Fas/Fas ligand system, especially systemic conseases associated with the Fas/Fas ligand system, especially systemic consease, Sjorgen's syndrome, pernicious or hypoplastic consenses, autoimmune hemolytic anemia, sterility, myasthenia gravis, condition, allergy, arteriosclerossis, myocarditis, cardiomyopathy, glomerulonephritis, hepatitis (fulninant, chronic, viral cells of the pative ligand, do not induce liver disease, and have actively mimic cells. They bind to both human and murine Fas, so can be evaluated in cells. They wind to both human and murine Fas, so can be evaluated in the native ligand, do not induce liver disease, and have reduced risk of inducing a human anti-murine antibody response. This sequence represents confirmer used in the construction of a humanised anti-Fas antibody HFE7A converted in the construction of a humanised anti-Fas antibody HFE7A converted in the method described in the construction of a humanised anti-Fas antibody HFE7A converted in the method described in the construction of a humanised anti-Fas antibody HFE7A converted in the method described in the construction of a humanised anti-Fas antibody HFE7A converted in the method described in the construction of a humanised anti-Fas antibody HFE7A converted to the construction of a humanise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New humanized anti-Fas antibody, useful for treating or preventing e.g. inflammatory or autoimmune disease, induces apoptosis selectively in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Serizawa N,
                                                                                                                                                                                                                                                                                                              30-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example reference 15; Page 137; 263pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cells with abnormal Fas-Fas ligand systems.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-258930/23.
                                             11-DEC-2001.
                                                                                                                                  Synthetic
                                                                                                                                                                                                  Human; mouse; Fas/Fas ligand system; Fas; antibody; light chain;
heavy chain; apoptosis; antiallergic; immunosuppressive; apoptot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SANY ) SANKYO CO LTD
28-MAR-2001; 2001JP-00093243.
                                                                                       JP2001342149-A
                                                                                                                                                                                                                                                                   Humanised anti-Fas antibody related
                                                                                                                                                                                                                                                                                                                                                                                                   ABL48723 standard; DNA; 20 BP
                                                                                                                                                                              autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7181 GGTGGGCATGTGTGA 7195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 GGTGGGCATGTGTGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                         disease; allergy; atopy; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haruyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     describes a novel humanized anti-Fas
                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nakahara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15; DB 1;
Pred. No. 1.9e+03
0; Mismatches 0
                                                                                                                                                                                                                                                                   PCR primer SEQ
                                                                                                                                                                                                immunosuppressive; apoptotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tamaki I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Takahashi
                                                                                                                                                                                                                                                                     ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody-like
                                                                                                                                                                                                                                                                     61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 2842
ABQ74807
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ঠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a preventive or treating agent for diseases caused by abnormality in the Fas/Fas ligand system containing, as the active component, an antibody having a light chain subunit and a heavy chain subunit and an activity of combining specifically with mammalian Fas and an activity of inducing apoptosis in a cell expressing Fas. The agent has antiallergic, immunosuppressive and apoptotic activity and is used for preventing and treating autoimmune diseases, allergy, atopy and others. The present sequence is that of a PCR primer useful in the construction of anti-Fas antibodies of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Drug for preventing or treating e.g. autoimmune disease or allergy, comprises humanized anti-Fas antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                         Tumour necrosis factor receptor 2; TNFR2; antisense oligonucleotide; phosphorothioate; 2'.-O-methoxyethyl; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 14 (preparatory); Page 32; 154pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-2000; 2000JP-00091144
                                                                         27-APR-2001; 2001US-00844634.
                                                                                                                           US6410324-B1
                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                        phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                              Human TNFR2 antisense oligonucleotide SEQ ID NO:57.
                                                                                                                                                                                                                                                                                                                                                                                                                       24-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                ABQ74807;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABQ74807 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SANY ) SANKYO CO LTD
                                                27-APR-2001;
                                                                                                    25-JUN-2002
                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7181 GGTGGGCATGTGTGA 7195
                                                                                                                                                                                                                                                                                                                                sapiens
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/mod_base= OTHER
/note= "2'-O-methoxyethyl nucleotides'
                                                                                                                                                                                           16. .20
                                                                                                                                                                                                                                                      /note= "phosphorothioate linkages"
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100.0%; Pr/
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Bennett CF,

Watt AT;

(ISIS-) ISIS PHARM INC

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RESULT 2843
ABL45980
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Best Local Similarity
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                       The invention relates to a preventive or treating agent for diseases caused by abnormality in Fas/Fas ligand system containing as the active component an antibody having as the light chain subunit a polypeptide containing residues 1-218 of one of 3, 239 residue amino acid sequences, or residues 1-451 of one of 3, 470 residue amino acid sequences, all fully defined in the specification and having an activity of combining specifically with mammalian Fas and an activity of inducing appytosis in a cell expressing Fas. The agent has immunosuppressive and antiallergic activity and is used for preventing and treating autoimmune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to a nucleic acid encoding human or mouse tumour necrosis factor receptor 2 (TNFR2). Also described is a method for inhibiting expression of human or mouse TNFR2 comprising contacting cells or tissues in vitro with one of the claimed compounds. The antisense compounds are used to treat a disease or condition associated with expression of TNFR2. The present sequence represents a human TNFR2 antisense chimeric phosphorothicate oligonucleotide, which is given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Humanised anti-Fas antibody related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compounds antisense to nucleic acid encoding human necrosis factor receptor 2 are useful to treat disease mouse tumor necrosis factor receptor 2 expression.
                                                                                                                                                                                                                                                                                                                                      29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; mouse;
light chain su
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-APR-2002
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                                                                                                                                                                                       Example 4 (Preparatory); Page 23; 194pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                     28-MAR-2001; 2001JP-00093106
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                                                                                                                                                                                                                                                                                                        (SANY ) SANKYO CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1765
                                                                                                                                                                                                                                      containing humanized anti-Fas antibody, used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chain subunit; apoptosis; immunosuppressive; antiallergic;
mmune disease; allergy; atopic; PCR primer; ss.
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                                                                                                                                                                                                                      autoimmune diseases, allergy, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   humanised anti-Fas antibody; Fas/Fas ligand;
               and others.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
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                                                                                                                                                                                                                                      preventing
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Best Local S
Matches 15
         The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC preventing a respiratory, lung or malignant disease or condition, also CC entiinflammatory steroid in a subject, for reducing or depleting levels of antiinflammatory steroid in a subject, for reducing or depleting levels of contiinflammatory producing bronchodilation, increasing levels of adenosine ceceptor, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or condition, inflammation, lung allergies, or a respiratory disease or condition. CC lung inflammation, lung allergies, or a respiratory disease or condition. The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Miller
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RESULT 2845
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CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or CC preventing a respiratory, lung or malignant disease or condition, also CC of, or reducing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels CC of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, CC Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at figure or a feeling in the printed specification, but was obtained in electronic format directly from WIPO
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     872pp; English
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ADA66464
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; antirheumatic; antiarthritic; gymecological; antiarteriosclerotic; Transforming Growth Factor beta-3; TGF beta-3; hyperproliferative disorder; cancers; atherosclerosis; rheumatoid arthritis; preeclampsia; fibrosis; phosphorothioate; ss.
                                    The present invention relates to antisense oligonucleotides (ADA66459-ADA66609), which inhibit Transforming Growth Factor (TGF) beta-3 expression. The oligonucleotides are useful for inhibiting the expression of TGF-beta3 in cells or tissues, and for treating an animal having a disease condition associated with TGF-beta3, e.g. a hyperproliferative disorder such as cancers of lung, liver, colon, osesphagus, pancreas, breast, skin or haematopoietic, atherosclerosis, rheumatoid arthritis,
                                                                                                                                                                 Novel antisense compound which is targeted to nucleic acid encoding transforming growth factor beta-3, and inhibits expression of TGF-beta 3, useful for treating a condition associated with TGF-beta 3, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA66464;
 Sequence
                         preeclampsia
                                                                                                                                          Claim 3; Page 87; 154pp; English.
                                                                                                                                                                                                                       WPI; 2003-229569/22.
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 BP;
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4 A;
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/note= "This oligonucleotide has a phosphorothioate backbone and 2-'methyoxyethyl (2'-MOE) wings at the and 3' ends, which are 5 nucleotides in length. Also cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                          fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
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 4 C; 4
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 T; 0 U; 0
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  Other;
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Query Match Best Local Matches 1

15;

Similarity

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Score 15; Pred. No.

DB 1; 1.9e+03

Length 20;

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Mismatches

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Gaps

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RESULT 2847
ABZ8096
XX
ABZ8096
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ABZ8096
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AC ABZ8096
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BERD97

RESULT 2848
ABZ80971
ID ABZ8097
XX ABZ8097
XX ABZ8097
XX ABZ8097
XX Primer;
XX Primer;
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FT modifie
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Best Local
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                 modified_base
                                                                  Human herpesvirus
                                                                                                                                                                     27-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 5; 6pp; Russian.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-146311/14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human herpesvirus.
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                                                                                                 Primer; PCR; diagnosis;
                                                                                                                                    Alternative
                                                                                                                                                                                                    ABZ80971;
                                                                                                                                                                                                                                   ABZ80971
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mmunobiology, virology and molecular biochemistry
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                                                                                                                                                                                                                                   standard;
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milarity 100.0%;
Conservative 0
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                                                                                                                                  Herpes virus infection diagnostic
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                               Location/Qualifiers
                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 C; 6 G; 5 T; 0 U; 0 Other;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
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                                                                                                                                  primer
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RESULT 2849
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Best Local (
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                                                                                                                                        Key
modified_base
                                                                                                                                                                                                                                                                                                                                                        Mouse; src-c; tyrosine kinase; src-c inhibitor; cytostatic; osteopathic; antiinflammatory; antibacterial; antisense therapy; vaccine; cancer; antisense oligonucleotide; aberrant bone remodeling; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ59521;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 two internal primers I and III (ABZ80969) on the second stage. The methe ensures determination of four types of herpes viruses in a single sample simultaneously. The method is useful in medicine, especially in immunobiology, virology and molecular biochemistry. This sequence represents an alternative version of primer III with an inosine residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method of diagnosing herpes virus infections by two-stage polymerase chain reaction (PCR) carried out using two external primers I (ABZ80967) and II (ABZ80968) on the first stage and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Method of diagnosis of herpes virus
                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                           phosphorothicate;
                                                                                                                                                                                                                                                                                                 Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                   hyperproliferative disorder; pancreatic cancer; lung cancer; tumour; ovarian cancer; oesophageal cancer; neuroblastoma; retinoblastoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse src-c chimeric phosphorothioate oligonucleotide SEQ ID NO:142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ59521 standard; DNA; 20 BP
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15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 3 A; 6 C; 6 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 3; 6pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001RU-00117331
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                                                                                                                                                                   Location/Qualifiers
                                                                                 /note=
/mod_base= OTHER
/note= "2'-O-methoxyethyl gapmer (2'-MOE wing)"
                                                                                                       /*tag= b
/mod_base=
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/note= "inosine"
                                                                                                                                                                                                                                                                           88.
                                                                                                                                                                                                                                                                                                 infection;
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                                                                                 "phosphorothicate linkages"
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                                                                                                       OTHER
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                                                                                                                                                                                                                                                                                                 inflammation;
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1.9e+03;
                                                                                                                                                                                                                                                                                                   tumour formation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a compound (I) that is 8-50 nucleobases CC in length targeted to a nucleic acid molecule encoding a 5'UTR, 3'UTR, CC coding region, intron region, exon region, stop codon, intron:exon CC punction, exon:exon junction, or 5' mRNA variant of src-c, and which CC specifically hybridises with and inhibits the expression of src-c. (I) CC have cytostatic, antiinflammatory, osteopathic and antibacterial CC activities, and can be used in antisense therapy and in vaccines. The CC entisense compounds (I) can be used for modulating the expression of src-c. c and for treating diseases or conditions associated with expression of CC src-c, e.g. aberrant bone remodeling or hyperproliferative disorders, CC garticularly cancer, such as breast cancer, neuroblastoma, retinoblastom CC cancer, overian cancer, oseophageal cancer, neuroblastoma, retinoblastom CC or Kaposi's sarcoma. (I) are also useful for diagnostics, therapeutics, CC prophylaxis, e.g. to prevent or delay infection, inflammation or tumour CC formation, as research reagents and kits, and in distinguishing between CC entertions of various members of a biological pathway. The present collegence represents a mouse src-c antisense chimeric phosphorothicate
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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useful for diagnosing, treating or preventing diseases associated wit
the expression of src-c, e.g. cancer or inflammation, and in research
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   Homo sapiens
                                                 Human; PR ramily remove , ...., PFM ZF domain; modulation of cell growth; cancer; PFM ZF domain; modulation of cell growth; cancer; cell degeneration disease; Alzheimer's disease; Parkinson's disease; insulin-dependent diabetes mellitus; IDDM; neuroprotective; insulin-dependent diabetes mellitus; IDDM; neuroprotective;
                                                                                                                                                                                                       06-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 16;
                                     reverse transcriptase-PCR; primer; ss.
                                                                                                                               Human; PR Family Member 5; PFM5; PFM PR domain; PFM zinc finger domain;
                                                                                                                                                                   Human PFM5
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                                                                                                                                                                                                                                                                               standard;
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                                                                                                                                                                   cDNA RT-PCR primer.
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llarity 100.0%;
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                                                                                                                                                                                                                                                                               DNA;
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== "2'-O-methoxyethyl gapmer (2'-MOE wing)"
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Pred. No.
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RESULT 2851
AAZ26500/c
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to the isolation of human and mouse PR Family Member (PFM) proteins, and the polynucleotide sequences encoding them. Also disclosed are PFM PR and PFM zinc finger (ZF) domains, and the polynucleotide sequences encoding them. The invention also discloses PFM cligonucleotides and methods for detecting a PFM polynucleotide sequence in a sample. The PFM polypeptide and polynucleotide sequences are useful for preparing a composition for modulating cell growth for treating cancer or diseases of cell degeneration, e.g. as Alzheimer's disease, parkinson's disease or insulin-dependent diabetes mellitus (IDDM). The present sequence represents a primer used in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New PR Family Member 2 oligonucleotide, useful for preparing a composition for modulating cell growth for treating cancer or discell degeneration, e.g., Alzheimer's disease or insulin-dependent diabetes mellitus.
                                                                                                                                                                      Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-SEP-1999;
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                                                                                                                                                                                                                                                                          Human polymorphic
                                                                                                                                                                                                                                                                                                        30-NOV-1999
                                                                                                                                                                                                                                                                                                                                                              AAZ26500 standard; DNA; 21 BP
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                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                    AAZ26500
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                                                                                                                                                                                                                                                                           region 689.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Lo. 1.9e+03;
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(VARI-) VARIAGENICS INC.

20-MAR-1997; 19-MAR-1998;

97US-0041057P

98WO-US005419.

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ARZ26584/c
ID AAZ2658
XX AAZ2658
XX AAZ2658
XX JO-NOV-
DE Human p
XX Polymor
KW cell vi
KW cell vi
KW allele
KW dysplas
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KW GERT
KW ALTHORO
OS HOMO SA
XX WO98416
XX YORN
PN WO98416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel method for identifying an inhibitor CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient having a CC precancerous condition, by administering to the patient a first allele CC specific inhibitor (ASI) trargeted to an allele of a first essential gene CC present in cells of the precancerous condition, where the normal somatic CC is active on at least one but less than all allelic forms of the gene CC present in a population and targets only one allelic form present in the CC used in the diagnosis, prevention and treatment of LOH disorders, e.g. CC cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic CC lesions, benign tumours, endometriosis, polycystic kidney disease, and CC malignant cells from bone marrow transplants. ARZ55812-Z26825 represent
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                                                                                                                                                                                                                                                                                                                                                                       Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metraplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
    WPI; 1998-521232/44
                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human polymorphic region 773.
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                                              Housman D,
                                                                                                                                         20-MAR-1997;
                                                                                                                                                                                                                                       24-SEP-1998
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                                                                                             VARIAGENICS INC
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                                              Ledley FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ledley
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                                                                                                                                            97US-0041057P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.2%;
                                              Stanton VP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stanton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ٧Þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
o. 2e+03;
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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밁 S

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used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic form present in the normal somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometriosis, polygetic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-C26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         \frac{\partial f}{\partial x}. Identifying target genes for allele-specific drugs – used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is ac
21
BP; 15
A;
0
C; 1
G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             active
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RESULT 2853
                                                       Matches
                                                                      Query Match
                                                                Local
                                    4464 TTTTTTTTTTTTT 4478
                       21
                                                      15; Conserv
                                                       Conservative
                                                              0.2%;
                                                      0,
                                                              Score 15;
Pred. No.
                                                       Mismatches
                                                              DB 1;
2e+03;
                                                                      Length 21;
                                                       Indels
                                                      <u>.</u>
                                                      Gaps
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0

밁 S

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AAZ74799/c
ID AAZ747
XX
AC AAZ747
AC AAZ747
AC Human
XX
Human
XX
Human
XW
Human
XW
Haploti
KW Haploti
KW diagno
XX
Homo s
XX
WO9954
XX
YX
Z1-APR
PR 21-APR
PR 21-APR
PR 21-APR
PR 21-APR
XX
XX
YX
YX
GEST
XX
XX
OGEST
XX
NOvel
PT Novel
PT Novel
PT Novel
PT Map of
                                                                                                                                                                                                                                               Human biallelic marker downstream amplification
                                                                                                                                                                                                                                                                                                     AAZ74799 standard; DNA;
                                                                                                                                                                                                                                                                  10-SEP-2001
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                              primer SEQ
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ID NO:9155.

Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer; Homo sapiens. diagnosis; ss

WO9954500-A2

28-OCT-1999.

21-APR-1998; 23-NOV-1998; 21-APR-1999; 98US-0082614P 98US-0109732P 99WO-IB000822

(GEST) GENSET

Cohen D, Blumenfeld 3 Chumakov

WPI; 2000-013267/01.

map of the human genome Novel biallelic markers used to construct a high density disequilibrium

Claim 8; Page 2182; 2745pp; English

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RESULT 2854
AAC73128/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
The present invention relates to an oligonuclectide array comprising oligonuclectide tags fixed to a solid substrate. The oligonuclectide array is useful for genotyping a nucleic acid sample at one or more larray is useful for genotyping a nucleic acid sample at one or more lavia single base extension (SBE) reactions. A pair of primers is used amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one such polymorphic locu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonuclectide array; genotyping; single base extension reaction; polymorphic locus; single nuclectide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ65654 to
                                                                                                                                 Universal array of oligonucleotides tags attached along with locus-specific tagged oligonucleotides using single base extension reactions.
                                                                                                                                                                                                                               Fan J,
Ryder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC73128 standard;
                                                                                                                                                                                                                                                                           (WHED )
                                                                                                                                                                                                                                                                                                                      26-MAR-1999;
23-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                               WO200058516÷A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-FEB-2001
                                                                                                        Example 7;
                                                                                                                                                                                                  WPI; 2000-656171/63
                                                                                                                                                                                                                                                                                                                                                                    27-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                 05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5151 GGGAGGGGAGTTCTC 5165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15;
                                                                                                                                                                                                                               Hirschhorn JN,
, Sklar P;
                                                                                                                                                                                                                                                                          WHITEHEAD INST
AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                        Page 49; 70pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ69578 represent human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                    2000WO-US008069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                      99US-0126473P.
99US-0140359P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #15 used
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                                                                                                                                                                                                                                                                                          BIOMEDICAL
                                                                                                                                                                                                                                             Huang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21
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                                                                                                        English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15;
Pred. No
                                                                                                                                                                                                                                             ×
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in multiplexing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                             Kaplan P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ŏ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ۲.
                                                                                                                                                                                                                                               Lander ES,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            markers from the present position 24 of their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR/SBE assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                     to a solid substrate useful in genotyping
                                                                                                                                                                                                                                               Lockhart DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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AAQ30432/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                             23-NOV-1990;
18-JAN-1991;
08-APR-1991;
17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     used in the present invention. The amplified nucleic acid product is used as a template in a SBE reaction with an extension primer. The seaction products are used to form the oligonucleotide array. Note: sequence includes a SNP represented by the degenerate codon in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21
                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                            Human interleukin-6 gene; herpes simplex; malignancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                     Oligomer IL6805
                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ30432 standard; DNA; 23
                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                           æ
                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                           malignancy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ30432;
                                                                                                                                                         25-NOV-1991;
                                                                                                                                                                         11-JUN-1992.
                                                                                                                                                                                          WO9209705-A1
                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                    misc_feature
                                                  Froehler B,
                                                                                             17-APR-1991;
17-APR-1991;
                                                                   (GILE-)
                                  1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 GGAGGCTGCGGGGGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                    GILEAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGAGGCTGCKGTGGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 4 A; 11 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                  Krawczyk
                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
                                                                                   91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
                                                                    SCI INC
                                                                                                                                        90US-00617907
                                                                                                                                                          91WO-US008811
                                                                                                                                                                                                                                                                                                                                                                                                     for forming triplex with HUMIL6 target duplex
                                                                                                                                                                                                                                                     /*tag=
/label=
                                                                                                                                                                                                          /mod_base= OTHER
/note= "OTHER= N
                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                             /note=
                                                                                                                                                                                                                                                                      /note= "o-xyloso
12. .23
                                                                                                                                                                                                                                                                                       *tag=
                                                                                                                                                                                                                                                                                                        note=
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                                                                                                                                                                                                                                                                                                                  base= OTHER
                                                  S
                                                                                                                                                                                                                                             "see
                                                                                                                                                                                                                                                                                                         "OTHER=
                                                                                                                                                                                                                                                      inverted
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                                                  Matteucci
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1;
                                                                                                                                                                                                                                            comments"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 2 T; 0 U; 1 Other;
                                                                                                                                                                                                             <u>4</u>
                                                                                                                                                                                                                                                                                                         Ŋ4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The amplified nucleic acid product is then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                     _polarity_region
                                                                                                                                                                                                                                                                               dimer synthon linkage"
                                                                                                                                                                                                                                                                                                         N4
                                                                                                                                                                                                             Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15;
                                                   Á
                                                                                                                                                                                                                                                                                                         ethanocytosine"
                                                                                                                                                                                                             ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
2e+03;
                                                  Milligan
                                                                                                                                                                                                                                                                                                                                                                                     AIDS; modified; HIV; RSV; HPV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.

which form a triplex with G-C and diagnosing HIV, hepatitis.

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RESULT 2856
AAQ03762
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The synthetic oligomer is capable of forming a triplex at physiological CC pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human CC interleukin 6 gene untranslated sequence contg. a purine rich sequence CC concd. on one strand of the duplex. The oligomer, and others like it are CC useful in diagnosis and therapy of diseases characterised by specific DNA CC duplex targets, e.g. HPV, HER, HTV, hepatitis B, herpes, malignant CC thmours and inflammation. The triple helices form under mild conditions CC thus assays may be carried out without subjecting the test specimen to thus assays may be carried out without subjecting the test specimen to CC harsh conditions. The oligomer contains an inverted polarity region CC formed from an o-xyloso dimer synthon. The linking gp. is o-xyloso (nucleotides have the 3'positions of xylose sugars linked via the o-CC xylene ring). Two nucleotides are coupled through a xylene residue to cligomer stable to nuclease activity. The oligomer is able to inhibit CC gene expression, as verified by in vitros systems. See also AAQ25452-25501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
Sequence containing this oligonucleotide encodes tPA-like polypeptides displaying the amino acid sequence of tPA in which 1-6 tripeptides are replaced by thr tripeptide RGD (=Arg-Gly-Asp). The produced polypeptides have plasminogen activator activity and can be used in fibrinolytic therapy, eg after cardiac infarct. They have increased blood clot specificity, longer half life, reduced inhibitor binding and/or greater proteolytic activity. See also AAQ03754-Q03771 and DE 3830734. (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue
                                                                                                                                                                                                                     WPI; 1990-100100/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
09-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Page
                                                                                                                                                                                   New tissue
                                                                                                                                                                                                                                                                                                                     17-SEP-1988;
                                                                                                                                                                                                                                                                                                                                                    17-SEP-1988;
                                                                                                                                                                                                                                                                                                                                                                                    22-MAR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                   DE3831714-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue plasminogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ03762;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ03762 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 0 A; 2 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                     Example 3.6;
                                                                                                                                                                                                                                                                                   (BADI )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4019 GAAAAAAGAGAGAAAACAAAATG 4041
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasminogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                  contg.
                                                                                                                                                                                                                                                  Schmidt M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                   plasminogen activator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                 Page 5; 17pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  71; 77pp; English.
                                                                                                                                                                                                                                                                                                                   88DE-03831714
                                                                                                                                                                                                                                                                                                                                                    88DE-03831714
                                                                                                                                                                  ninogen activator analogues - are polypeptide(s) with arginine-glycine-aspartic acid tri:peptide units.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    analogue oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tPA analogues; fibrinolytic therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                 Baldinger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 23;
                                                                                                                                                                                                                                                    Schwarz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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AAQ21598/c
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-AUG-2003
25-MAR-2003
16-MAY-1992
                                 -FMDV sequence and subjected to exonuclease II digestion to remove sequences encoding the 39 N- terminal amino acids from the IBRV gIII gene. The final plasmid obtained has the sequences for bGH-FMDV whih cabe expressed from the IBRV gIII promoter. When the Pst1 to Mlul region AAQ21592 is transferred into the pLA(GIII):30 dl HindIII/Hpal oligo 171/172 cassette, the bGH-FMDV sequence will be in the same reading frast the GIII protein. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                 insertion into the NCOI site of the IBRV gIII gene-contg. plasmid pLA(GIII):30 dl. Hind III/Hapl (see AAQ21591). The cassette was designed to provide cloning sites, optimal translation start signals for the bovine growth hormone (bGH) signal sequence, the RMDV epitope sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         on 25-MAR-2003 to correct PD field.)
                                                                                                                          and restriction nuclease sites to permit unidiretional nulease digestion into the IBRV gene. The resulting plasmid was then transformed with a bGH
                                                                                                                                                                                                                                                                                 Vaccine
                                                                                                                                                                                                                                                                                                          WPI; 1992-058515/08.
                                                                                                                                                                                                                                                                                                                                   Dimarchi RD,
                                                                                                                                                                                                                                                                                                                                                           (NOVA-)
                                                                                                                                                                                                                                                                                                                                                                                                  24-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                          22-JUL-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-FEB-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP471457-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bovine herpesvirus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Infectious bovine rhinotracheitis virus primer oligo EJ48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ21598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ21598 standard;
                                                                                                                                                                                                       An oligonucleotide cloning/expression
                                                                                                                                                                                                                              Example; Page 63; 72pp; English.
                                                                                                                                                                                                                                                                     virus-based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Foot-and-mouth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Herpes virus-based viral vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3524 GACCCTGTCTTCCTTCCGCCCGC 3546
                      cassette, the bGH-FMDV sequence will be gIII protein. (Updated on 25-MAR-2003 to don 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                           NOVAGENE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                     viral vector
                                                                                                                                                                                                                                                                               foot and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                  Kit S,
                                                                                                                                                                                                                                                                                                                                                                                                  90US-00556593
                                                                                                                                                                                                                                                                                                                                                                                                                           91EP-00306646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                             ELI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                                                                                 mouth
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                                                                                                                                                                                                                                                                                                                                   Little
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23
                                                                                                                                                                                                                                                                     ch disease and
which expresse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 4 T; 0
                                                                                                                                                                                                                                                                     expresses
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    foot
                                                                                                                                                                                                                                                                                                                                   Gale
                                                                                                                                                                                                      cassette was synthesised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and mouth disease; epitope;
                                                                                                                                                                                                                                                                     es a foot and mouth disease virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           踞
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                                                                                                                                                                                                                                                                                                                                   Ki t
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Best Local Similarity

0.2%;

Score 15; Pred. No.

2.2e+03;

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23;

Query Match

Sequence

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                                                                                                                                                                                                                                                     RESULT 2859
AAT40331/c
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                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
               Wild type; self-splicing group I intron; large ribosomal RNA precursor; Tetrahymena thermophila; catalysis; enzymatic RNA; food product; anti-viral agent; mutation; personal care product; cleaning agent; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chromosomal DNA from Thermosipho africanus (Taf) was PCR-amplified with degenerate primers corresponding to the amino acid sequences of conserved regions of known thermostable polymerases. DG129 is one example of a degenerate primer. See AAQ23917 for full-length Taf Pol I coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thermostable DNA polymerase from Thermosipho africanus - prepd. purificn. from cells or by expression of Taf polymerase gene in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thermostability; polymerase chain reaction; Taf Pol Thermosipho africanus; thermophilic bacteria; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Degenerate sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ23949 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-150887/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9206202-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-OCT-1992
                                                                                                                                             06-DEC-1996
                                                                                                                                                                                                                                 AAT40331 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1992.
                                                                                                   DNA cleavage
                                                                                                                                                                                       AAT40331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CETU ) CETUS
                                                                                                                                                                                                                                                                                                                                                                                   4687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7234 CCTCTCAAGTCCAGCATGGATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                           15;
                                                                                                                                                                                                                                                                                                                                         4 GATCTGGNGAYGAYCCSATG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Page 38; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
                                                                                                                                                                                                                                                                                                                                                                               GATCTGGTGATGAAGCCATG 4706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCTCTCAAGTTGGGCACGGAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer DG129 based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 5 A; 4 C; 7 G; 3 T; 0 U; 4 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lawyer FC,
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                                                                                                 substrate for generation of
                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORP.
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                                                                                                                                                                                                                                 명
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15; DB 1;
Pred. No. 2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Greenfield L,
                                                                                                   improved ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                             ω
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Matches
                                                                                              Query Match
                                                                                                                                                                                          The sequences given in AAT40331-32 represent sequences that were as substrate molecules in experiments for selection of improved catalytic activity of ribozymes. The evolution experiment spanned 10 successive generations and catalytic activity was deduced after each generation. The self-splicing group I intron of the invention is based on the large ribosomal RNA precursor from Tetrahymena thermophila. The biological function of this molecule is to catalyse its own excision from precursor rRNA to produce mature rRNA. The Tetrahymena wild type sequence was used in the design of the enzymatic RNA molecules of the invention. A number of mutations are listed in the specification which improve the enzymatic properties of this molecule, e.g. G444A, G191U, U190A and A314G. The modified enzymatic may be used as medical or pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                           Enzymatic RNA molecules having one or more point mutation(s) - the enzymatic performance of the molecules.
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01-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                Sequence 23 BP;
                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 111; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-010936/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-NOV-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W09531551-A1
                                                                                                                                                                              agents for use in anti-viral agents, food products, personal care
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SCRI ) SCRIPPS
                                                                                ocal
                  6683 TATTTTTATTTATATATGGGGCC 6705
23
                                                                                Similarity
 TTTATTTATTTAGAGGGCC
                                                                 Conservative
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94US-00270180
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                                                                                                                                12 A; 4 C; 2 G; 5 T; 0 U; 0 Other;
                                                                              0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INST
                                                                                                                                                                agents
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                                                                                Score 15; DB 1;
Pred. No. 2.2e+0;
                                                                  Mismatches
                                                                                .20+03;
                                                                                                 Length 23;
                                                                  Indels
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RESULT 2860 AAT40328/c wild type; self-splicing group I intron; large ribosomal RNA precursor; Tetrahymena thermophila; catalysis; enzymatic RNA; food product; anti-viral agent; mutation; personal care product; cleaning agent; ss. Synthetic. Group I intron substrate 05-DEC-1996 AAT40328; AAT40328 standard; WO9531551-A1 (first entry) DNA; 23

Gaps

0

13-MAY-1994; 01-JUL-1994;

94US-00242402 94US-00270180 95WO-US005141

26-APR-1995; 23-NOV-1995.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequences given in AAT40327-30 represent sequences that were used to CC optimise DNA cleavage activity of the enzymatic RNA molecule of the CC invention. The 3' potion of the substrate was transferred to the 3' CC terminal G of the ribozyme and amplification was performed. The product CC of the reaction was a molecule which contained the 3' portion of the CC substrate attached to the 3' end of the ribozyme. Selection occured when CC a primer was hybridised across the ligation junction and used to initiate CC and thus led to selective amplification of the catalytically active CC RNA's. The self-splicing group I intron of the catalytically active CC arge ribosomal RNA precursor from Tetrahymena thermophila. The CC large ribosomal RNA precursor from Tetrahymena thermophila. The CC large ribosomal RNA produce mature rRNA. The Tetrahymena wild type sequence CC was used in the design of the enzymatic RNA molecules of the invention. A CC number of mutations are listed in the specification which improve the CC enzymatic properties of this molecule, e.g. G44AA, G191U, U190A and CC natally agents for use in anti-viral agents, food products, creament are readucts for use in anti-viral agents, food products,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Sim
Matches 18;
                                                                                                                                                                                                                                                                                                  Primer; intron-5; Arabidopsis; RPP5; disease-resistance; RT-PCR; reverse transcription; nested polymerase chain reaction; downy mildew; Peronospora parasitica; transgenic plant; crop improvement; ss.
                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT37491 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enzymatic RNA molecules having one or more point mutation(s) - the enzymatic performance of the molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-010936/01.
Isolated Arabidopsis pathogen resistance gene RPPS - for prodn.
                                WPI; 1996-465029/46.
                                                                 Jones
                                                                                                                                   07-APR-1995;
                                                                                                                                                                    09-APR-1996;
                                                                                                                                                                                                     10-OCT-1996.
                                                                                                                                                                                                                                    WO9631608-A1
                                                                                                                                                                                                                                                                                                                                                                    Intron-5 primer for RPP5 transcript RT-PCR analysis
                                                                                                                                                                                                                                                                                                                                                                                                       19-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 BP; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       personal care products or cleaning agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1;
                                                                                                  (INNE-) INNES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SCRI ) SCRIPPS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TATTTTTATTTATATATGGGGCC 6705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTATTTATTTAGAGGGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 97; 209pp; English.
                                                                 Parker J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                  CENT INNOVATIONS LTD
                                                                                                                                   95GB-00007232.
                                                                                                                                                                    96WO-GB000849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 4 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                 Coleman M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15; DB
Pred. No. 2.2e
0; Mismatches
                                                                 Daniels MJ,
                                                                                                  JOHN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                 Szabo
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This primer may be used with primers AAT37483-90 and AAT37492 in RT-PCR analysis of a gene (AAT37476) encoding Arabidopsis RPP5 protein, which conferrs disease-resistance against the downy mildew fungus (Peronospora parasitica). First strand cDNA synthesis from seedling leaf mRNA and PCR amplification have been performed using intron flanking primers. Primers AAT37489-90 are used in the first amplification in a nested PCR strategy for identification of intron-5, and this primer and primer AAT37492 are used in the second amplification. Cloned PCR products have been sequenced using vector-specific and insert-specific primers, to identify introns and exons. Primers corresponding to conserved regions in the protein may be used to identify other resistance genes in plants. The RPP5 gene may
                                                                                                                                                                                                                                                                                                                                                                                                                                              Asthma;
chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                     Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of
                                                                                                                                                                                                                              WPI; 1997-051871/05
                                                                                                                                                                                                                                                                                                            07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                         W09640162-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23 BP; 5 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transgenic plants esp. resistant to downy mildew fungus
                                                                                                                                                                                                                                                                                                                                      06-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                19-DEC-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Al adenosine receptor antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT76031 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        be expressed in a transgenic plant to confer disease-resistance
                                                                                                                                                                                                                                                                                   (UYEC-)
                                                                                                                                                                                                                                                      JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 TCGCAGTGGGAGCTGCTGCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      н
                                                                                                                                                                                                                                                                                   AIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                obstructive
                                                                                                                                                                                                                                                                                                                                                                                                                                                            airway epithelium; adenosine free; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TCTCAATGTGAGCGGCTGCAAGC
                                                                                                                                                                                                                                                      Metzger WJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 33; 59pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                   EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                            95US-00474497
                                                                                                                                                                                                                                                                                                                                       96WO-US009306
                                                                                                                                                                                                                                                                                                                                                                                                                                              pulmonary disease; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15; I
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23
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which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide specific for the human AI adenosine receptor. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON, its liberation upon antisense degradation is prevented, thereby

A method for treating airway disease in a subject has been produced

Example 5; Page 23; 71pp; English.

preventing

adenosine-induced bronchoconstriction in

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SXS
                                                                                                                                                                   RESULT 2863
AAV37972/c
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RESULT 2864
AAV09049/c
ID AAV09049 standard; DNA; 23 BP
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                                                               В
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Best Local :
                                                                                                                 Best Local Similarity
Matches 18; Conserv
                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 0 A;
                                                                                                                                                                                         The present sequence represents a construction oligonucleotide for ECEPO section 2 as part of the assembly of human exythropoietin (EPO), used in an example from the present invention. The present invention describes recombinant human EPO which causes bone marrow cells to increase production of reticulocytes or red blood cells, where the polypeptide is the product of expression in CHO (Chinese hamster ovary) cells of an exogenous DNA sequence encoding human EPO. EPO is potentially useful in the diagnosis and treatment of blood disorders characterised by low or defective red blood cell production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reactive airways
                                                                                                                                                                                                                                                                                                                                               Recombinant human erythropoietin - potentially useful for diagnosis and treatment of blood disorders.
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-261957/24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; erythropoietin; EPO; bone marrow; reticulocyte; red blood cell;
expression; CHO; chinese hamster ovary cell; diagnosis; blood disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ECEPO section 2 construction oligonucleotide 9 for human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV37972;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV37972 standard; DNA;
                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-FEB-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AU688723-B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                       Example 11; Page 64; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                       (KIRI ) KIRIN AMGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5699
                                                                                       7124
                                                               23
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18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTGCCTTCCTTTTCCTCTTCTC 5721
                                                                              TTCCTGTGCACACAGTCCAGCCT 7146
                                                                                                                                                                  23 BP; 4 A; 6 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTTTCCTTCCTTTGTCTCTCTTC 23
                                                                 TGCCTGGCCACGCAGTACAGCCT 1
                                                                                                                 0.2%;
llarity 78.3%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97AU-00046867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97AU-00046867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8 C; 1 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23
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                                                                                                                 0; Mismatches
                                                                                                                             Score 15;
Pred. No.
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Pred. No. 2
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                                                                                                                               N
                                                                                                                                          DB 1;
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5;
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                                                                                                                 5
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                                                                                                                                          Length 23;
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RESULT 2865
AAT99526
ID AAT9952
XX
AC AAT9952
XX
DT 21-MAYXX
DE Human S

AAT99526 standard; DNA;

23

0

21-MAY-1998

(first entry)

Human ST receptor PCR primer.

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                                                                                                                                                                               This sequence is a substrate for a wild type tetrahymena ribozyme L-21 CC form. The ribozyme sequence is an example of a catalytic RNA (I) of the CC invention, which catalyses site-specific cleavage of nucleic acid under CC physiological conditions includes a sequence derived from a group I CC intron. Similar catalytic RNAs (II) which catalyse hydrolysis of amide CC ends are useful as peptidases and proteases, e.g. in wound debridement, CC clot dissolution, in detergents or as a meat tenderiser. (I) cleave CC single- and (partly) double-stranded nucleic acids in vitro or in vivo, CC and are potentially useful as antiviral agents and gene regulators; also CC to generate defective but still immunogenic viruses (for vaccines); CC diagnostically to detect mutations in nucleic acid or to identify nucleic CC acid binding agents; to modulate/terminate reactions initiated by DNA CC therapeutic/diagnostic processes using antisense sequences; in DNA CC therapeutic/diagnostic processes that provide better catalytic performance; CC in vitro evolution processes that provide better catalytic performance; CC consideriation in temperature and pH ranges; new enzymatic activities or constitutions.
                                                                    Matches
                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Catalytic RNA for site-specific cleavage of nucleic acid or hydrolysis amide bonds - and ribozyme amidase intermediates, useful e.g. as peptidase(s), antiviral agents and gene regulators.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mutation detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tetrahymena ribozyme; group I intron; amide end hydrolysis; peptidase; protease; antiviral agent; gene regulator; immunogenic virus; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Substrate for tetrahymena ribozyme L-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-110627/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Joyce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV09049;
                                                                                                                                   Sequence 23
                                                                                                                                                                      specificities; altered recognition sites or co-factor requirement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SCRI ) SCRIPPS RES INST
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                   6683 TATTTTTATTTATATATGGGGCC 6705
23
                                                                     18;
TTTATTTATTTAGAGGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 120; 215pp;
                                                                                                                                    BP;
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                      12 A; 4 C; 2 G; 5 T; 0 U; 0 Other;
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                                                                                  0.2%;
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                                                                                    Score 15;
Pred. No. 2
                                                                     Mismatches
                                                                                                        DB 1;
                                                                                    2e+03
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                                                                     Indels
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AAX34877
ID AAX3487
XX AAX3487
AC AAX3487
AC AAX3487
XX ZB-JUN-
XX JE PCR Pri
XX Immorta
XX Immorta
XX SV40 vi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claimed PCR primers (see AAT99462-T99531) hybridise to sequences that encode the extracellular domain of human heat-stable toxin (ST) receptor protein (see AAM33371), a highly specific marker for metastasised colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in AAT99526 and AAT99527. Claimed in vitro methods for determining whether or not (i) an individual has metastasised colorectal cancer cells, or (ii) a tumour cell is a colorectal cancer cell comprise the steps of examining a sample of extraintestinal tissue and/or body fluids or tumour cells from an individual to determine whether ST receptor protein is being expressed by cells in the sample. Expression is determined by immunoassay or by PCR using primers that selectively amplify ST receptor cDNA or mRNA (see also
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           using primers that AAT97229)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Determining whether an individual has metastasised colorectal cancer cells and origin of tumour cells - by detecting presence of heat-stal toxin receptor on cells in a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Waldman SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ST receptor; heat stable toxin receptor; colorectal metastasis; diagnosis; human; PCR; primer; ss.
                06-APR-1999
                                             JP11089565-A
                                                                                                                  Immortalized human hair papila cell; HPC; screening; SV40 viral Large T-antigen gene; deleted replication
                                                                                                                                                              PCR primer used
                                                                                                                                                                                           28-JUN-1999
                                                                                                                                                                                                                        AAX34877;
                                                                                                                                                                                                                                                  AAX34877
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYJE-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                        3199
                                                                                                    growth
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                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
18; Conserv
                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                       AGTGAGGGGCTTGAGAAAGTGGG 3221
                                                                                                                                                                                                                                                                                                                                                                                                                                               23 BP;
                                                                                                                                                                                                                                                                                                                           AATGAGGGGCTGGAAATAGTGAG 23
                                                                                                    stimulating
                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JEFFERSON THOMAS
                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Carrithers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-0016564P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97WO-US007467.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              8 A; 1 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                             amplify FGF4.
                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
78.3%;
                                                                                                    agent; PCR primer;
                                                                                                                                                                                                                                                    23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SL;
                                                                                                                                                                                                                                                    ВР
                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                   Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detecting presence of heat-stable
                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                   .2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                               Length 23,
                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
                                                                                                                  hair growth; initiation point;
                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumour;
                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                          impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain, cystic fibrosi; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; panoreatic cancer; colon cancer; breast cancer; lung cancer; panoreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The specification describes the preparation of immortalized human hair papila cells (HPC). The method comprises transformation of HPC with an SV40 viral large T-antigen gene with deleted replication initiation point. The immortalized HPC can be used in a screening method for a hair growth agent, by culture of immortalized HPC in the presence of a substance to be tested and observation of the growth of the immortalized HPC. HPC is also used in development of hair growth stimulating agents. The resent sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the course of the sequence represents a PCR primer, which is used in the sequence of the sequence represents a PCR primer.
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                                                                                                                                          17-SEP-1997;
09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                           colon cancer; breast cancer; lun
hepatocellular carcinoma; kidney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human adenosine Al receptor antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 23 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-SEP-1997;
                                                                                                                                                                                                                                                  25-MAR-1999
                                                                                                                                                                                                                                                                                         WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                          prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX53824 standard; DNA;
                                                                                                   (UYEC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SHIS ) SHISEIDO CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         374 ACTACGAGGTGGACATCAAGCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
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                                                                                                     VIND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACTACCTGCTGGGCATCAAGCGG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                     EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                          97US-0059160P.
98US-00093972.
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                                                                                                                                                                                                         98WO-US019419
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                                                                                                                                                                                                                                                                                                                                                                            88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15; DB
Pred. No. 2.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                             g cancer; m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                               hepatic metastasis;
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WPI; 1999-229400/19

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RESULT 2868
AAX18455/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those essociated with impaired respiration and inflammation, including lung CC diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, CC acute asthma, allergies, asthma, impeded respiration, respiratory clistress syndrome, pain, cystic fibrosis, pulmonary hypertension, CC pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary conditions are those such as leukemias, lymphomas, carcinomas e.g. Cc colon cancer, breast cancer, lung cancer, menanoma, hepatic metastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                         Graves' disease; thyroid adenoma; hypertension; cardiomyopathy; schizophrenia; Kaposi's sarcoma; fibroblast growth factor receptor; adenylate cyclase constitutive activator; thyrotropin receptor; thyrotropin stimulating hormone; beta-adrenergic receptor; PCR primuthyrotropin stimulating hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX18455;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX18455 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotides used in treatment of, e.g. pulmonary
                                                                                                                                                                                     14-APR-1998;
                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                             Agonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer for DNA encoding orphan homologue GPR12.
                                 WPI; 1999-105468/09
                                                                   Behan DP,
                                                                                                                                                     14-APR-1997;
                                                                                                                                                                                                                                                         WO9846995-A1
                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                       22-OCT-1998
                                                                                                   (BEHA/) BEHAN D P.
(CHAL/) CHALMERS D T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              identification; orphan receptor; constitutively active OR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTTGCCTTCCTTTTCCTCTTCTC 5721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Trirectricerrierererere 23
                                                                   Chalmers DT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 42; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                     97US-00839449
                                                                                                                                                                                       98WO-US007496.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8 C; 1 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    В₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15; DB
Pred. No. 2.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                             PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coding and
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identification of agents that reduce receptor activation, rather than compounds that antagonise the normal ligand. Once identified, (inverse) agonists can be used to study OR function. The method does not require knowledge of the endogenous receptor ligand or receptor function, and identifies directly compounds that inhibit the activated receptor, i.e. able to block both ligand-dependent and -independent activation, rather than only the ligand-dependent process, as is the case with compounds identified by ligand-dependent assays. It should accelerate drug discovery at a wide range of OR and since activated receptors have a greater response to the agents, potential drugs are more likely to be
                                                                                                                                                                                                                                                                                                                                                                              GPR3. The invention relates to a method for the identification of candidate compounds as agonists, including inverse or partial, of an orphan receptor (OR), which comprises: (i) applying test compound to constitutively active OR; and (ii) measuring its effect on OR. The method is particularly used to identify inverse agonists of G protein-coupled OR, i.e. potential therapeutic agents for treating conditions in which constitutively active OR are implicated (e.g. Graves disease, thyroid adenoma, hypertension, cardiomyopathy, schizophrenia, major depression, Kaposi's sarcoma and many others tabulated). It is based on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             constitutively active receptor - particularly therapeutically active inverse agonists at G protein coupled receptors, without requiring knowledge of endogenous ligand or receptor function.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence is a primer for DNA encoding the oirphan receptor homologue GPR3. The invention relates to a method for the identification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72; 114pp; English.
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Best Loc Matches Query Match Local 1669 CAACCTTGTTTCTGCAAATATGC 1691 23 18; Similarity CACCCATGTTCCTGCTAATAGGC Conservative 0.2%; 0, Score 15; Pred. Mismatches ŏ. 2e+03; 5 Length 23; Indels 0 Gaps

0,

Sequence 23

B₽;

δ

A; 4 C; 8 G; 5 T; 0 U; 0 Other;

RESULT 2869 Human Y chromosome-specific sequence DYZ1 PCR primer YZ1. 16-MAR-1999 AAX00055; AAX00055 standard; DNA; 23 (first entry)

hybridisation; kidney barrier; foetal sex; maternal; inherited disease; cancer; pathogenic infection; paternity testing; PCR primer; ss. Human; Y chromosome; DYZ1; urine; detection; amplification; diagnosis;

Synthetic Homo sapiens

WO9854364-A1

03-DEC-1998. 29-MAY-1998; 98WO-US010965

30-MAY-1997; 03-JUN-1997; 97US-0048170P. 97US-0048381P.

(LXRB-) LXR BIOTECHNOLOGY INC

٨, Melkonyan HS, Umansky

WPI; 1999-070223/06

Identifying agonists of orphan receptors from their effect on

Analysis of DNA present in urine that has crossed the kidney barrier useful in the determination of foetal sex, and in the diagnosis of

inherited

disease,

cancer and pathogenic infections

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RESULT 2870
AAA3326
XX
AAA3326
XX
AC AAA3326
XX
DE Low ad
XX
Human;
KW phosph
KW allerg
KW antial
KW lung (
KW respi:
KW respi:
KW cance
XX
OS Homo
XX
PP 03-4
XX
PR 03-4
XX
PR 03-7
XX
XX
PR 03-7
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XX
PR 03-7

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; altergic disease; bronchoconstriction; inhibitor; antinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hyperiension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A method has been developed for the analysis of a fragment of foetal DN that has crossed the placental and kidney barriers. The method comprise assaying for presence of foetal DNA in a urine sample from a pregnant subject. Also described are: (1) detecting any target nucleic acid sequences in a urine sample; (2) diagnosis for male foetal DNA in maternal urine; and (3) general method for detecting Y Chromosomespecific nucleic acids using the primers given in AAX00055 to AAX00058. The methods are used: (1) to determine the sex of a foetus (by detectin respecific fragments); (11) for diagnosing or monitoring diseases,
                                                                                                                                                                                                                 New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-FEB-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Low adenosine antisense oligonucleotide SEQ ID NO:956.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23 BP; 2
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                                                                                                                  English
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Pred. No. 2
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CC nucleic acids involved in bronchoconstriction, allergies, and/or CC inflammation. The ON can have antiinflammatory, antiallergic, CC inflammation. The ON can have antiinflammatory, antiallergic, antiiasthmatic, cytostatic and analgesic activities. The compositions are CC useful for the treatment of diseases associated with inflammation, CC impaired airways, including lung disease and diseases whose secondary CC effects afflict the lungs of a subject. They can be used for treating CC e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic CC fibrosis, pulmonary hypertension, emphysema, chronic obstructive CC pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, CC carcinomas, and cancers which may metastasise to the lungs, including CC breast and prostate cancer. The reduction of the adenosine content of the CC onstruction and inflammation. AAA32313 to AAA3312 represent the CC nucleotide sequences given in the sequence listing from the present content of invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 equences are also called SEQ ID NO:1 to 185, but the sequences differ CC CAAA3392) are specifically claimed ONs from the present invention on N.B. Sequences given in the disclosure of the present invention do not match CC up with their corresponding SEQ ID NO: sequences given in the sequence given in the sequence of the present invention of the sequence content of the content of the content of the present invention of the sequence of the present invention of the sequence content of the content
Sequence 23
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RESULT 2871 AAA64522/c AAA64522 standard; DNA; 23

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PCR primer IntABR used to amplify exon 1 of human FEZ1 gene

02-JAN-2001

(first entry)

Human; FEZ1 gene; tumour suppressor gene; 8p22; cancer; tumour growth; tumour proliferation; tubulin; microtubule; protein EF1-gamma; tubulin polymerisation disorder; mitosis initiation; cell proliferation; cell growth; cell shape; cell rigidity; cell motility; DNA replication; tumorigenesis; survival; metastasis; PCR primer;

Homo sapiens

WO200050565-A2

25-FEB-2000; 2000WO-US004950

25-FEB-1999; 99US-0121537P

(UYJE-) UNIV JEFFERSON THOMAS

Croce ã Ishii H;

WPI; 2000-558396/51

New polynucleotide homologous with a portion of one strand of the human FEZ1 gene, useful for alleviating abnormal cell proliferation such as

Example 1; Page 45; 255pp; English

The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets

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ARESULT 2872
ARAZ50566/c
ID ARZ50566/c
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Matches
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          The patent discloses a method of identifying agonists and inverse or partial agonists to the endogenous, constitutively activated G protect coupled orphan receptors (GPCRs), by contacting them with a GPCR fus: protein comprising a GPCR and a G protein. Determining expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G protein-coupled orphan receptor; GPCR; agonist; G protein;
GPCR fusion protein; inverse agonist; drug; treatment; PCR p
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                                                                                                                                                                                                             identifying a compound
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ilarity 78.3%;
Conservative
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constitutively activated G protein-coup
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09-JUN-1998;
09-JUN-1998;
(II) is an adenosine AZa receptor agonist (IA), or an oligonuclectide (IB), containing less than 15% adenosine (A), that is antisense to target genes or corresponding RNA, to genomic flanking regions (i.e. 5' or 3' ends or segments between coding and non-coding sequences), or to all segments of mRNA encoding the adenosine A1, A2a, A2b or A3 receptors, and has A1, A2b or A3 agonist activity or A2a antagonist activity (or at least no agonist activity at this receptor). (I) may be a mixture of (Ia) and (Jb), and optionally also contains one or more surfactants. The compositions are used to prevent, alleviate and/or treat adenosine receptor-mediated cardiac, lung and/or renal damage or failure (particularly where associated with ischaemia, toxin release and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           endotoxin release; ARDS; acute respiratory distress syndrome; cytoprotective; anti-allergic; anti-inflammatory; anti-hypoxic; supraventricular tachycardia; allergic rhinitis; acute inflammation; chronic obstructive pulmonary disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; adenosine Al receptor; antisense oligonucleotide; adenosine Ala receptor; adenosine Ab receptor; adenosine phosphorothicate; cardiopulmonary failure; renal failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              agonists to these receptors can be used as drugs for treating related diseases. The present sequence is a PCR primer used 220 bp fragment of human GPRI2 (G protein-coupled receptor) examine tissue samples for expression of these receptors
                                                                                                                                                                                                                                                                                                                                                           Nyce
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA03669 standard; DNA; 23
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                                                                                                                                                                The present invention describes a pharmaceutical composition, con at least one agent (I) that prevents, alleviates and/or inhibits adenosine-mediated cardiopulmonary and/or renal damage and/or fa:
(I) is an adenosine A2a receptor agonist (Ia), or an oliconucleoi
                                                                                                                                                                                                                                              Claim 17; Page 38; 252pp;
                                                                                                                                                                                                                                                                                               Novel
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                                                                                                                                                                                                                                                                                          composition for treating or preventing e.g. cardiopulmonary and
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                                                                                                                                                                                                                                                                                                                                                           Hill JL;
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98US-00093972.
98US-0088657P.
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nypoxia;
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RESULT 2874
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                               Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-679539/66.
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nilarity 78.3%;
Conservative
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14
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cc expression and or activity of target polypeptides associated with cc lung/respiratory disorders and malignancies, such as stimulating and citivating peptide factors and transmitters, transcription factors, cimmunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, communoglobulins and antibodies, antibody receptors, cytokines and commonkine receptors, adenosine receptors, binding proteins, adenosine receptors, brokens expected common receptors, cytokine and compressed system (CNS) and peripheral nervous and non-nervous system peptide transmitters, defensins, growth factors, vascative peptides and creceptors, binding proteins and malignancy associated proteins. The contisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or lung inflammation, allergy(ies) and/or and/or lung inflammation, allergy(ies) and/or condition selected from pulmonary vasoconstriction, inflammation, condition selected from pulmonary vasoconstriction, bronchitis, condition selected from pulmonary vasoconstriction, bronchitis, condition selected from pulmonary vasoconstriction, pulmonary infections, bronchitis, condition cancer. AAF18434 to AAF21543 represent human polynucleotide condition of the present invention.
Sequence 23
                                                                        present invention
BP; 0 A; 8 C; 1 G;
14 T; 0
U; 0 Other;
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Matches Query Match Best Local : 5699 18; ۳ Similarity Trirectricerrierererere 23 rrrecerrecerrrecerrere 5721 Conservative 0.2%; 0 Score 15; I Pred. No. 2 Mismatches DB 1; 2.2e+03; 5; Indels Length 0 Gaps

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AAA39003 standard; DNA;
   23
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30-AUG-2000 (first entry)

Human orphan receptor GPR12 RT-PCR primer #2

Identification; modulator; cell surface membrane receptor; treatment; orphan receptor; antithyroid; antidiabetic; neuroleptic; antidepressant; cytostatic; G protein-coupled receptor agonist; reverse transcriptase; primer; 88

Homo sapiens.

WO200021987-A2

20-APR-2000.

12-OCT-1999; 99WO-US023935

13-OCT-1998; 98US-00170496

(AREN-) ARENA PHARM

WPI; 2000-317935/27.

Identifying compounds with inverse agonist activity to orphan receptors useful for treating e.g. Graves' disease, and schizophrenia, involves contacting candidate compounds with constitutively activated receptors.

4. Page 73; 110pp; English

The present invention describes a method for directly identifying

The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities.

Claim 14; Page 120; 1592pp; English

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RESULT 2876
AAD19181/c
ID AAD1918
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Best Local
27-APR-2000;
27-APR-2000;
14-AUG-2000;
18-DEC-2000;
18-DEC-2000;
22-DEC-2000;
28-DEC-2000;
28-DEC-2000;
04-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         agonist activity to a constitutively active orphan receptor (ORR). The method comprises determining the efficacy of the compound by contacting it with the ORR. A compound identified by the above method having inverse agonist activity to ORR is useful for the treatment of diseases characterised by constitutive activation of the receptor e.g. Graves' disease, male precocious puberty, Jansen's disease, retinitis pigmentosa, hypoparathyroidism, neuropsychiatric diseases, schizophrenia, major depression, and cancerous growth in Kaposi's sarcoma. The method can identify (I) directly without prior knowledge or use of receptor ligands and is useful for accelerating drug discovery at a broad range of ORR. The present sequence represents a reverse transcriptase PCR primer for the human orphan receptor GFR12, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; G-protein coupled receptor 12; GPCR12; cardiomyopathy; vaccine; atherosclerosis; diabetes; cardiant; cytostatic; cancer; obesity; pain; diabetes mellitus; anorexia; cachexia; cardiomyopathy; atherosclerosis; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; immune disorder; haematopoietic disorders; retinal disorder; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                              human immunodeficiency virus; adenocarcinoma; bulimia; asthma; ulcer; angina pectoris; hypotension; hypertension; Crohn's disease; anxiety; multiple sclerosis; schizophrenia; dementia; mental retardation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human GPCR12
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                                                                                                                                                      27-APR-2001;
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                                                         2000US-0199947P.
2000US-0199960P.
2000US-0275226P.
2000US-0256399P.
2000US-0256524P.
2000US-0258159P.
2000US-0258511P.
20COUS-0258828P.
2001US-0259659P.
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                                                                                                                                                       2001WO-US013680
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                                                                                                                                                                                                                                                                                                                                                                                                                                 osteoporosis; urinary retention;
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/mod_ba
                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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/note= "FAM-labelled adenosine"
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                                                                                                                                                                                                                                                 "TAMRA-labelled thymidine'
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Pred. No. 2.2e+03;
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RESULT 2877
AAD19184/c
ID AAD19184 standard; DNA; 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present DNA sequence is a probe which is used for determining the CC cNA. GPCR protein and DNA may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate GPCRX expression, CC obesity, diabetes mellitus, anorexia, cachexia, cardiomyopathy, pain, CC atherosclerosis, neurodegenerative disorders (Alzheimer's disease, Runtington's disease); bulimia, immune disorder, CC parkinson's disease, Huntington's disease); bulimia, immune disorder, CC naematopoietic disorders (Bioorders (Bioorders), CC catheroscopietic disorders, disorders related to cell signal processing and CC metabolic pathway modulation, retinal disorder (photoreception), CC bacterial, fungal, protozoal and viral infections (HIV); cancer (neoplasm CC cychn's disease, multiple sclerosis, hypotension, hypertension, asthma, CC (dementia, mental retardation, schizophrenia, anxiety); acute heart CC failure, osteoporosis, myocardial infarction and urinary retention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                               Human; G-protein coupled receptor 12; GPCR12; cardiomyopathy; vaccine; atherosclerosis; diabetes; cardiant; cytostatic; cancer; obesity; pain; diabetes mellitus; anorexia; cachexia; cardiomyopathy; atherosclerosis; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; immune disorder; haematopoietic disorders; retinal disorder; HIV; human immunodeficiency virus; adenocarcinoma; bulimia; asthma; ulcer; angina pectoris; hypotension; hypotension; lorder; disease; anxiety; multiple sclerosis; schizophrenia; dementia; mental retardation; gene therapy; osteoporosis; urinary retention; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G-Protein coupled receptor polypeptides and NAs useful for preventing diagnosing and treating cardiomyopathy, atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Padigaru M, I
Alsobrook JP,
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                                                                                                                                                                                                                                                                                                                                                                   Human GPCR12 cDNA expression analysis determining probe Ag1609.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 23 BP; 6 A; 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Macdougall JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-MAR-2001; 2001US-00275226
                                             modified_base
                                                                                                               modified_base
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                                                                                                                                                                  sapiens
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Burgess CE, Casman
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                                                /note=
                                                                                                                                  Location/Qualifiers
               /*tag= b
/mod_base= OTHER
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/mod_base= OTHER
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5
                                                                "FAM-labelled adenosine"
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0; Mismatches
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SJ, Lepley DM, Gango
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Gangolli
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/note= "TAMRA-labelled thymidine"

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ID AAH4041
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Best Local
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27-APR-2000;
14-AUG-2000;
Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       parkinson's disease, Huntington's disease); bulimia, immune disorder, haematopoietic disorders, disorders related to cell signal processing and metabolic pathway modulation, retinal disorder (photoreception). descrerial, fungal, protozoal and viral infections (HIV); cancer (neoplasm adenocarcinoma); angina pectoris, hypotension, hyperrension, asthma, Crohn's disease, multiple sclerosis, ulcers, neurological disorders (dementia, mental retardation, schizophrenia, anxiety); acute heart failure, osteoporosis, myocardial infarction and urinary retention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present DNA sequence is a probe which is used for determining the expression analysis of human G-protein coupled receptor-12 (GPCR-12) cDNA. GPCR protein and DNA may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate GPCRX expression, obesity, diabetes mellitus, anorexia, cachexia, cardiomyopathy, pain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   obesity, diabetes mellitus, anorexia, cachexia, cardiomyopathy, pain, atherosclerosis, neurodegenerative disorders (Alzheimer's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G-Protein coupled receptor preventing, diagnosing and t
                                                                                                                                                          SNP specific lower
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28-DEC-2000;
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18-DEC-2000;
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15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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2000US-0256524P.
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2001US-0259659P.
2001US-00275226.
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2000US-0275226P.
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Burgess CE, Cas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6 A;
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                                                                                                                                                          PCR primer SEQ ID 3206.
                                                                                                                                                                                                                                                                                                                   DNA;
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Pred. No.
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SJ, Lepley DM, Gango
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                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
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Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                               inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific for a human SNP containing DNA sequence
                                                                                                                                            AAF32064;
                                                                                                                                                                            AAF32064 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
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Pred. No.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tomato, Arabidopsis and carnation cDNA clones encoding senescence-induced deoxyhypusine synthase and eIF-5a, useful for inhibiting senescence in a plant when introduced in reverse orientation into the genome of the
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19-JUN-2000; 2000US-00597771.
                                                                                       02-JAN-2001.
                                                                                                                                                                                                                                                                                  MEK kinase 1; MAP/ERK kinase kinase 1; pro-apoptotic; apoptosis signal regulation; programmed cell death; serine/threonine kinase; MAP kinase cascade; JNK/SAPK; substrate; Jun N-terminal kinase/stress-activated protein kinase; Bcl-2 substrate; NF-kappa-B-mediated transcription regulation; expression inhibition; antisense therapy; hyperproliferative disorder; cancer; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                         Human MEKK1; mitogen-activated protein kinase kinase kinase MEK kinase 1; MAP/ERK kinase kinase 1; pro-apoptotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human MEKK1
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                                23-JUL-1999;
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                                99US-00359756.
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Pred. No. 2.26
0; Mismatches
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RESULT 2881
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound targeting nucleic acid encoding human mitogenactivated protein kinase kinase 1 (MEKK1), useful for treating diseases or conditions associated with MEKK1 expression, or preventing inflammation or tumor formation.
                                                                                                                                                                                                          Human; ss; G protein coupled receptor; GPCR; insulin secretion; diabetes; antidiabetic; pancreatic beta cell; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                          diabetes; antidiabetic; pancreatic intracellular cAMP concentration.
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                                               30-NOV-2001; 2001WO-JP010472
                                                                                                                       WO200244362-A1
                                                                                                                                                        Homo sapiens
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ilarity 78.3%;
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01-DEC-2000; 2000JP-00367349 10-AUG-2001; 2001JP-00243841

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tomato; ss; PCR; deoxyhypusine synthase; DHS; senescence; eIF-5A; primer; eukaryotic initiation factor 5A; plant; cell death; disease resistance; antisense; blossom end rot; environmental stress; pathogen resistance; shelf-life; perishable fruit; flower; vegetable.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     agents using G-protein coupled receptor (GPCR) polypeptide which promotes insulin secretion by pancreas beta-cells under high glucose levels, and increases intracellular cyclic adenosine monophosphate (cAMP concentration. Also included are (1) screening method of cells transformed by a vector containing a gene encoding the GPCR polypeptide; (2) antidiabetic agents identified by the screening method; (3) drug compositions containing these compounds; and (4) treatment of diabetes using these drug compositions. The present sequence is a 5 RACE (rapid amplification of cDNA ends) PCR primer used to isolate the human cDNA amplification of cDNA ends) PCR primer used to isolate the human cDNA
                                      Increasing resistance to physiological disease in plant, by integrating gene or its fragment encoding senescence-induced deoxyhypusine synthase or eIF5A in antisense orientation into plant genome and growing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Drug screening using a polypeptide which promotes pancreatic beta cell insulin secretion and intracellular cyclic adenosine monophosphate concentration for identification of antidiabetic agents.
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Ueda Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
               16-OCT-2000;
26-OCT-2000;
31-OCT-2000;
03-NOV-2000;
12-DEC-2000;
17-DEC-2000;
22-JAN-2001;
22-JAN-2001;
                                                                                                                                                                                                                                                                                    G protein coupled receptor; GPCR; olfactory receptor; cell signal processing disorder; metabolic pathway modulation; cardiomyopathy; atherosclerosis; diabetes; developmental disease; immune disease; taste disorder; scent detectability disorder; obesity; burkitt's lymphoma; corticoneurogenic disease; infectious disease; pain; signal transduction pathway disorder; metabolic pathway disorder; retinal disease; metabolic disorder; cancer; parkinson's disease; acute heart failure; urinary retention, osteoporosis; Crohn's disease; ulcer; allergy; neurological disorder; genetic disorder; transplantation; fertility; Pancreatitis; Hyperthyroidism; Endometriosis; forensic biology; transgenic animal; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; Page 44; 114pp; English
                                                                                                                                                                                            23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel G-protein coupled receptor probe #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK95525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                           16-OCT-2001;
                                                                                                                                                                                                                            WO200240539-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK95525 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCTTTCTCTCTAGGATTCTTT
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ilarity 78.3%;
Conservative
2000US-0240704P.
2000US-0244497P.
2000US-0244548P.
2000US-0255017P.
2000US-0263216P.
2001US-0263216P.
2001US-0263118P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B₽;
                                                                                                                                                              2001WO-US032256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    relates to increasing resistance to physiological disease
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>و</u>
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ARBSULT 2884
AAD32230/c
ID AAD3223
XX AAD3222
XX AAD3222
XX
DT 18-JUN-
XX
DE RASSF1/
XX Nested
KW neck;
KW antiser
KW antiser
KW gene tl
XX
OS Uniden
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC useful for identifying an agent that binds to the polypeptides are cceptor. GPCRX polypeptides are confectively receptor and date substance or ligand molecules interacting with an colfactory receptor polypeptide. The polypeptide, [1] and [1] are also consistent for treating diseases and disorders related to cell signal consistent diseases, taste and scent developmental diseases, immune cc diseases, taste and scent detectability disorders, Burkitt's lymphoma, cc corticomeurogenic disease, signal transduction pathway disorders, cc corticomeurogenic disease, signal transduction pathway disorders, cc corticomeurogenic disease, signal transduction pathway disorders, cc diseases, taste and scent detectability disorders, Burkitt's lymphoma, cc diseases, transplantation, cancer, parkinson's disease, acute cheart failure, urinary retention, osteoporosis, Crohn's disease, ulcers, cc allergies, neurological disorders, genetic disorders, transplantation, cf trility, pancreatitis, Hyperthyroldism and Endometriosis. GPCRX csequences are also useful for identifying acell or tissue type in a biological sample, to amplify DNA sequences from very small biological samples such as tissues e.g. hair or skin or body fluids in forensic control of continuing and/or activity of GPCRX protein and for identifying and/or evaluating modulators of GPCRX protein and for identifying and/or evaluating modulators of GPCRX protein and for detect DNA encoding a convert grant compled receptor produced from real time quantative (RTQ) cc -PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESERVATION OF THE STATE OF THE
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Best Local
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Peyman JA,
Lepley DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-FEB-2001; 2001US-0268225P.
15-FEB-2001; 2001US-0269031P.
27-JUL-2001; 2001US-0308203P.
                                                                                                          Nested polymerase chain reaction; gene inactivation; cancer; lung; head; neck; leukaemia; colorectal; prostate; bladder; immunological modulation; antisense treatment; alkylating treatment; aberrant methylation; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel G protein coupled receptor especially olfactory receptor polypeptides and nucleic acids for diagnosing and treating atherosclerosis, cardiomyopathy and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Colman SD,
                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23 BP; 6
                                                                                  gene therapy;
                                                                                                                                                                                                                RASSF1A gene amplifying reverse unmethylated PCR primer.
                                                                                                                                                                                                                                                                       18-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                            AAD32230 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 223; 309pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 267 GCAGGTGTTCCAGGC 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 100.0%;
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Ballinger RA, Padigaru M, Wolenc AR, Shenoy SG;
Gerlach V, Gangolli BA, Macdougall JR, Smithson G;
Stone DJ, Gunther E, Ellerman K, Grosse WM, Alsobrook JP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Burgess CE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 23;
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AR, Shenoy SG;
TR, Smithson G;
Alsobre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Best Local :
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CC (PCR) for amplifying a gene that is altered in a particular cancer. The content of comprises expanding copies of a gene by amplification method and CC using the product in a second methylation-specific amplification to detect gene inactivation. The method is useful for amplifying the gene CC cancers which include head and neck, leukaemia, colorectal, prostate and CC bladder) thereby permitting cancer detection and monitoring by detecting GC gene inactivation in biological fluid such as sputum and blood. The method finds application in the area of chemoprevention, where markers GC are needed to identify high-risk subjects and to evaluate efficacy of CC preventive agents. The method is further useful for identifying cincer this premailgnant state, and for monitoring the efficacy of the interventions, by determining whether the previously detected methylation CC biomarker persists or disappears during the course of the interventions, which include immunological modulation, antisense treatment, gene therapy CC and alkylating treatments. The method is useful for detecting the CC aberrant methylguanine DNA methyltransferase gene (MGMT), RASSFIA) gene or other gene promoters. The present sequence is CC acceptable or amplifying RASSFIA gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nested, two-stage polymerase chain reaction for amplifying gene in cancer e.g. of lung, comprises expanding copies of gene by amplification method and using the product in a second amplification to detect gene
    Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Belinsky SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-AUG-2000; 2000US-0228057P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-AUG-2001; 2001WO-US026452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a nested, two-stage polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (LOVE-) LOVELACE RESPIRATORY RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200218649-A1
B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21; 55pp; English.
    14 A; 8 C; 0 G; 1 T; 0 U; 0 Other;
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Matches 3167 GTTAGGTTTGGGTTTGATACTTT 3189 Conservative 0, Pred. No. ... 0; Mismatches Indels

Similarity

0.2%;

Score 15; DB 1; Pred. No. 2.2e+03;

DB 1;

Length 23

0

Gaps

0

23

GTTTGGTTTGTGTTAGTGT 1

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RESULT 2885
ABL99466/c
ID ABL9946
XX ABL9946
XX ABL9946
XX CABL9946
XX CADLE
CY CALL
C
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Left PCR primer used to target TGFB-1 canine gene ABL99466; 02-JUL-2002 (first entry

Canine gene array; toxicological response;

WO200208453-A2

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Best Local S
Matches 18
                                                                                                                                                                                                     progressive systemic sclerosis; recurrent pericarditis;
Sjogren's syndrome; relapsing phychondritis; arthritis; rheumatism;
sarcoidossis; SLE; splenic atrophy; diabetes; Wegener granulomatosis;
ulcerative colitis; vasculitis; vitiligo; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to identifying a toxicologically relevant canine gene and the generation of an array of toxicologically relevant canine genes. The gene array is useful for obtaining a gene expression profile, by exposing a population of cells to an agent, obtaining cDNA from the population of cells, labeling the cDNA, and contacting the cDNA with the gene array. The relevant gene is useful for making and using arrays to determine toxicological responses to various agents, and also useful for identifying novel gene sequences and novel canine genes. The method for analysing toxicological responses using the canine gene array is rapid and efficient. The present sequence is related to the canine gene array
                                                                                                                                                                                                                                                                   Al thyroid disease; atrophic gastritis; pernicious anaemia; Chron's disease; colltis ulcerosa; Goodpasture syndrome; IgA nephropathy; IgG glomerulonephritis; myasthenia gravis; partial lipodystrophy; polymyositis; primary biliary cirrhosis; primary sclerosing cholangitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying toxicologically relevant canine gene to determine toxicological responses of agents, by obtaining and comparing gene expression profiles of untreated canine cells and canine cells treated canine cells are cannot can be considered.
                                                                                                                                                                                                                                                                                                                                   Transglutaminase; TGM; transamidation; autoimmune disease; Addison's disease; AI haemolytic anaemia; AI thrombocytopenic purpara;
                                                                                                                                                                                                                                                                                                                                                                                    Novel transglutaminase TGz associated PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK67730 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 5; Page 53; 140pp; English.
                              15-SEP-2000;
16-MAY-2001;
                                                                                                                                            WO200222830-A2
                                                                                                                                                                                                                                                                                                                                                                                                                       02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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(UYCA-) UNIV COLLEGE CARDIFF
                                                                             14-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4735
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 Similarity
18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHASE-1 MOLECULAR TOXICOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCCAGCTGGAGGAAGAAGGGTC 4757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGCCATGAGGAGCAGGAAGGGTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                              2000GB-00022768.
2001GB-00011995.
                                                                             2001WO-GB004120.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 C; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.2e+03;
5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 23;
                                                                                                                                                                                                                                                                                                                                                                                     #17.
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The invention relates to nucleic acids which encode novel polypeptides CC having transglutaminase activity. The compositions of polypeptide are useful for transamidation reactions on peptides and polypeptides. CC useful for transamidation reactions on peptides and polypeptides. CC in a diagnostic method in a subject or in cells derived from a subject CC having an autoimmune disease. The method for detecting transglutaminase proteins may be used to diagnose autoimmune diseases which include proteins may be used to diagnose autoimmune diseases which include cc hyroid disease, AI haemolytic anaemia, AI thrombocytopenic purpara, AI convoid diseases, atrophic gastritis, pernicious anaemia, Chron's CC disease, colitis ulcerosa, Goodpasture syndrome, Igh nephropathy or IgG colymoratitis, primary biliary cirrhosis, primary sclerosing cholangitis, cpolymositis, primary biliary cirrhosis, primary sclerosing cholangitis, cpolymositis, rheumatoid arthritis, rheumatism, sarcoidosis, Sjogren's syndrome, SLE, splenic atrophy, type I (insulin-dependent) diabetes complities, Wegener granulomatosis, ulcerative colitis, vascuiltis (both systemic and cutaneous) and vitiligo. This sequence represents a primer cused in the study of transglutaminase genes in which DNA, amino acid constances and chromosomal locations of novel transglutaminases are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acids which encode novel transglutaminase enzymes TG-Z and TG-Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aeschlimann DP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Grenard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnostic methods
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diseases.
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밁 ঠ Matches Query Match Best Local : 4731 TGGAGGCCAGCTGGAGGAAGAAG 18; -Similarity TĠAĀĠCTĊĀĠĊCĠĠĀĠĠTĀĠĀĀĠ 23 Conservative 0.2%; 0, Score 15; Pred. No. Mismatches 4753 DB 1; 2.2e+03; 5 Length Indels °, Gaps

0

Sequence 23

BP; 7

Α,

4 C; 9

e G

3 T;

0 U;

0 Other;

ABZ95083; ABZ95083 standard; DNA; (first entry) 23

Human adenosine Al receptor antisense fragment no.946.

antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung al Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; inflammation; respiratory disease; gb allergy;

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM

Nyce JW, Miller S, Li Y, Tang Sandrasagra A, L, Shahabuddin S Katz Ħ Pabalan Ģ Aguilar D;

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RESULT 2888
ABX93429
ID ABX9342
XX
AC ABX9342
XX
DT 27-MAY
XX Chro
XX
DE Y Chro
XX
UTransg
KW Iransg
KW Specii
KW specii
KW specii
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
              30-MAY-1997;
03-JUN-1997;
29-MAY-1998;
04-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or corresponding transfer of the corresp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid detection; nucleic acid modification detection; specific foetal nucleic acid; cancer; diabetes; arteriosclerosis; obesity; autoimmune disease; chromosomal abnormality; genetic disease; haemophilia; Alzheimer's disease; Huntington's disease; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Y chromosome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pathogen infection; genetic predisposition; cancer treatment monitoring; foetus sex determination; foetal genetic disease; paternity; amniocentesis; Y chromosome specific sequence; DYZ1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAY-2003
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                                                                                                                                                                                                                                         03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                  10-DEC-2002
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nilarity 78.3%;
Conservative
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              97US-0058170P.
97US-0048381P.
98WO-US010965.
2000US-00230704.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        material monitoring; urine sample analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specific sequence DYZ1,
                                                                                                                                                                                                                                         2000US-00634732.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        kidney barrier;
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misc_feature

/mod_base= '/note= "5'

polylinker"

*tag=

modified_base

Location/Qualifiers

88

Synthetic. microarray;

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The invention describes a method of monitoring (M) transplanted material containing providing a urine sample suspected of containing nucleic acid of the transplanted material, where the transplanted containing containing the urine sample for nucleic acids from the cell genome of the transplanted contained contained that are different from nucleic acids of the transplanted contained that that are different from nucleic acids of the transplanted contained that the transplanted nucleic acids as well as nucleic acid modifications and contain composed the kidney barrier. (M) is useful for detecting the presence of specific nucleic acids as well as nucleic acid modifications and contain composed the transplanted contained nucleotides, for diagnosis of diseases such as cancer, diabetes, arteriosclerosis, beesity and various autoimmune diseases or chromosomal containing that transplanted contains and pathogen infections, for containing contains to various diseases, for monitoring cancer treatment, for analysing specific nucleic acids in urine to track the success of transplanted cells, tissue and organs, for determination of foetus sex and the identification of foetal genetic diseases, such as those inherited from the father for various purposes, including containing DNA modifications accompanied by death of at least one subset of the cells bearing DNA modification. (M) is a safe, and simple non-invasive test, an excellent alternative to capacific DYZ1 sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 18; Conserv
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74137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monitoring transplanted material in a patient, by analyzing urine samples for nucleic acids from cell genomes of the transplanted material that are different from nucleic acids of recipient and have crossed the kidney
                                                                                                                                                                 G-protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-340405/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-JUL-2000; 2000US-00609162
                                                                                                                                                                                                     Oligonucleotide used
                                                                                                                                                                                                                                      23-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23
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                                                                                                                                                                                                                                                                         ABV74137
                                                                                                                                                                                                                                                                                                       ABV74137 standard; DNA; 23 BP
                                                                                                                                                                                                                                                                                                                                                                                                                           5692 CCACTGTTTGCCTTCCTTTTCC 5714
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                                                                                                                                                  coupled receptor; odourant; receptor; olfaction;
y; anosmia; attractant; arometic; pesticide; PCR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
                                                                                                                                                                                                      in cDNA library array
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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Gaps

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                                                                                                                                                  The present sequence is that of an oligonucleotide used in a method CC designed to rapidly array and normalize a complex cDNA library obtained CC from a target species. Clones are arrayed into multi-well plates. Each CC well contains 16 oligonucleotides with a 5' polylinker, a poly-T run CC which allows an anchored oligonucleotide in each well to selectively CC which allows an anchored oligonucleotide in each well to selectively CC unique 3' key sequences are designed to give a comprehensive level of CC degeneracy since they are diverse and numerous enough to ensure that CC every possible cDNA sequence can be bound by an individual, specific CC oligonucleotide in a single well. The cDNA library is heated to denature CC the clones into single stranded DNA, and an aliquot is added to every common 5' region present in every cDNA clone serves as the 5' priming CC well. The anchored oligonucleotide serves as the 3' primer in PCR, and CC site. Denaturing and washing leave anchored cDNA in each well. The CC and isolate clones encoding 6-protein coupled receptors, especially CC pathway of invertebrates and vertebrates, e.g. odourant binding proteins, CC or other olfactory or neuronal proteins. The identified receptors and crive effectors involved in the olfactory capanimal's sensitivity to odours, for manufacturing compounds or devices that mask odours, or trapping invertebrates with odourants.

CC semicochemicals (e.g. arometics or pheromone mimetics) can be developed controlling insect feeding and breeding behaviour, detecting the presence of small air-borne molecules, etc
                                                               Best Local
Matches
                                                                            Query Match
Best Local :
                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying nucleic acid encoding novel sex-linked-tissue-linked receptors, useful for isolating odorant binding proteins or pesticide alternatives, by analyzing sequences from a male- and female-specific nucleic acid library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 5; 83pp; English.
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31-JAN-2002;
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22. 23
/*tag= C
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ABZ21707 standard; DNA; 23 BP

片 S

5953 CAAGCTTATACTAGAGAAGAAGCA 5975

1 CATGCTTCTACTGAAGAACAGCA 23

Query Match Best Local

18;

Conservative

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Mismatches

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Similarity

0.2%;

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DB 1; 2.2e+03; 0 Other;

Length 23;

Sequence

23 BP;

æ P.

6 C; 4 G; 5 T;

0 U;

RESULT 2890

ABZ21707

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The present invention describes an RNA-binding motif protein-15 (RBM15)-CC megakaryoblastic leukaemia-1 (MKL1) fusion protein, a MKL1-RBM15-S fusion protein, a mKL1-RBM15-S fusion protein, and a MKL1-RBM15-S+AE fusion protein associated with acute proteins associated leukaemia (AMKL). Also described: (1) an antibody that specifically binds to the RBM15-MKL1, MKL1-RBM15-S, or MKL1-RBM15-S+AE CC fusion protein; (2) a non-human transgenic animal that has been altered to express a gene encoding a RBM15-MKL1, MKL1-RBM15-S, or MKL1-RBM15-S+AE CC fusion protein; (3) identifying an agent capable of binding to a RBM15-CC MKL1, MKL1-RBM15-S, or MKL1-RBM15-S+AE fusion protein; (4) detecting the t(1;22) chromosomal rearrangement associated with AMKL; and (5) screening CC protein arising from the t(1;22) chromosomal rearrangement associated with AMKL; and (5) screening CC with AMKL. The fusion proteins have cytostatic activity of a fusion CC proteins and nucleotide molecules are useful for designing and preparing agents that specifically inhibit the expression of the RBM15-S+AE fusion CC proteins and nucleotide molecules are useful for designing and preparing CC asymmet that specifically inhibit the expression of the RBM15-MKL1 or MKL1-RBM15 genes in cells for therapeutic and other purposes. The transgenic CC canimals are useful for identifying and testing carcinogenic or therapeutic compositions. The methods are also useful for attenting the t(1;22) chromosomal rearrangement associated with AMKL, or for identifying agents useful for treating patients with AMKL. The antibodies CC can be used to selectively, kill cells expressing RBM15-MKL1, MKL1-RBM15-S+AE fusion protein. RBM15 is located to chromosome 1 pagents and preparity of a fusion of the present sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New RNA-binding motif protein-15 (RBM15)-megakaryoblastic leukemia-1 (MKL1), MKL1-RBM15-S and MKL1-RBM15-S+AE fusion proteins, useful for identifying agents useful for treating patients with acute megakaryoblastic leukemia.
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                                                 represents a PCR primer which is used in the generation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RBM15-MKL1 fusion protein PCR
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in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Page 108; 109pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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cytostatic;
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RESULT 2891

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                                                                                                                                                                           CC deposited at the UK National Collection of Type Cultures under number CC 11740. Also described: (1) selecting an agent for treating a Mycoplasma CC infection in a subject by assaying a sample from the subject for CC Mycoplasma spp A39, and if the assay is positive, selecting as an agent CC enalogue or derivative; (2) an oligonucleotide probe which is capable of Mybridising to a region of polynucleotide from Mycoplasma spp A39 and if the assay is positive, selecting as an agent CC corresponding region from another Mycoplasma; (3) an oligonucleotide probe which is capable of Mybridising to a polynucleotide or corresponding region from another Mycoplasma; (3) an oligonucleotide or corresponding region from another Mycoplasma; (3) an oligonucleotide corresponding region from another Mycoplasma; (3) an oligonucleotide corresponding for Mycoplasma spp A39 and incapable of amplifying a sequence cof the same length on the target sequence from another Mycoplasma; and CC (4) assaying for Mycoplasma spp A39 and incapable of amplifying a sequence cof the same length on the target sequence from another Mycoplasma; and CC (4) assaying for Mycoplasma spp A39 by obtaining a sample potentially containing a sample potentially containing a sample potentially containing the length of the 16S-23S (bp). The Mycoplasma spp A39 or its component can be used as a control in an assay of a sample potentially containing Mycoplasma spp A30. Compounds Compositions for the treatment of a human subject with Mycoplasma spp A39. Compounds composition for the subject is capable of production of an antibiotic composition of an assay for Mycoplasma spp A39 or its component, and the CC compound comprises a pleuromatalin consisting of valuemalin or its compound comprises a pleuromatalin consisting of valuemalin or its component or or derivative. The present sequence represents a PCR primer for the 16S rDNA region of Mycoplasma spp A39, which is given in the
                                                              Query Match
Best Local
                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated Mycoplasma spp A39 (UK NCTC number 11740), useful control in assaying samples potentially containing Mycoplasma and for selecting agents against Mycoplasma infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JUL-2000; 2000GB-00017073.
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                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 10; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (WIND/) WINDSOR H.
                                          18;
                                                              Similarity
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                                                                                                                                                                   the present invention
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1 spp A39,
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ABZ59392;
                       ABZ59392 standard; DNA; 23
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RRESULT 28
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XX IS 17-APR-2003 Mouse src-c probe (first entry) SEQ ID NO:13

Mouse; src-c; tyrosine kinase; src-c inhibitor; cytostatic; osteopathic antiinflammatory; antibacterial; antisense therapy; vaccine; cancer; antisense oligonucleotide; aberrant bone remodeling; breast cancer; hyperproliferative disorder; pancreatic cancer; lung cancer; tumour; ovarian cancer; oesophageal cancer; neuroblastoma; retinoblastoma; Kaposi's sarcoma; infection; inflammation; tumour formation; probe; ss. Mus musculus osteopathic;

WO200295053-A2

28-NOV-2002.

16-MAY-2002; 2002WO-US015684.

18-MAY-2001; 2001US-00860473

ISIS PHARM INC

2003-120806/11.

New antisense oligonucleotides targeted to nucleic acids encoding src-c, useful for diagnosing, treating or preventing diseases associated with the expression of src-c, e.g. cancer or inflammation, and in research

Example 13; Page 85; 137pp; English

The present invention describes a compound (I) that is 8-50 nucleobases CC in length targeted to a nucleic acid molecule encoding a 5'UTR, 3'UTR, CC coding region, intron region, exon region, stop codon, intron:exon function, exon:exon junction, or 5' mRNA variant of src-c, and which cc specifically hybridises with and inhibits the expression of src-c. (I) cc have cytostatic, antiinflammatory, osteopathic and antibacterial cc artivities, and can be used in antisense therapy and in vaccines. The cc antisense compounds (I) can be used for modulating the expression of src-c c and for treating diseases or conditions associated with expression of CC c and for treating diseases or conditions associated with expression of CC src-c, e.g. aberrant bone remodeling or hyperproliferative disorders, CC particularly cancer, such as breast cancer, neuroblastoma, retinoblastoma cancer, ovarian cancer, oesophageal cancer, neuroblastoma, retinoblastoma cor Kaposi's sarcoma. (I) are also useful for diagnostics, therapeutics, Crophylaxis, e.g. to prevent or delay infection, inflammation or tumour continuous of various members of a biological pathway. The present continuous represents a probe for mouse src-c, which is used in an example of from the present invention. sequence represents a probe from the present invention

Sequence 23 ₿P; 4 Ð, 7 C; 7 <u>ი</u> v ij 0 U; 0 Other;

S 밁 Matches Best Local Query Match 30 Similarity GAGCTGCTGCAGGCT 44 Conservative 0.2%; Score 15; pred. No. 0 Mismatches DB 1; Lo. 2.2e+03; Length 23; Indels <u>.</u> Gaps

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CC nucleotide sequence encoding peroxisome proliferator-activated receptor-cC diabetes of at least one nucleotide, or comprising a fragment of the nucleotide of at least one nucleotide, or comprising a fragment of the concluded of the concluded of the nucleotide of a transparent of the concluded of the nucleotide of a mutation in the gene encoding concluded of the concluded of the nucleotide of the concluded of the nucleotide of the concluded of the nucleotide of the second labelled polynucleotide to the nucleotide of the nucleotide of the substance with which the second labelled polynucleotide to the nucleotide of the nucleotide of
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AAL55157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel mutant DNA sequence encoding peroxisome proliferator-activated receptor coactivator useful for identifying subjects who are at increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peroxisome proliferator-activated receptor-gamma coactivator-1; PGC-1; type 2 diabetes; antidiabetic; enzyme; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated polynucleotide molecule comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUN-2001;
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7022 CAGAGGAMAATAGGAMACCCTCC 7044

Query Match Best Local Matches

18;

Similarity

0.2%;

Score 15; DB Pred. No. 2.2e 0; Mismatches

2e+03;

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Gaps

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RESULT 2895 ABV77669/c

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Length 23; Indels

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Sequence

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BP; 13

A; 6 C; 3

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ABX79809/c
                                                                                                                                                       repeat within a coding sequence (expressed sequence tag, EST), which comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep.X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABT95676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying a candidate polymorphic repeat within a coding sequence, understanding or treating genetic disease, comprises detecting tande repeats in a target coding sequence and scoring the repeats for polymorphic probability.
                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                       The invention discloses a method for identifying a candidate polymorphic
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                              4017 GAGAAAAAAGAGAGAAAACAAAA 4039
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                                                                              Similarity
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                                                                                                                            1 C; 0 G; 23 T; 0 U; 0 Other;
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                                                                               Score 15;
Pred. No.
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RESULT 2896
ABA98547/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; zinc fit
anti-HIV; PCR;
                               14-JUN-2000; 2000CN-00116509.
                                                                          14-FEB-2002
                                                                                                                                     Insulin-like growth factor; binding protein; cytostatic;
cancer; PCR primer; ss.
                                                                                                                                                                     Insulin-like
                                                                                                                                                                                        30-APR-2002
                                                                                                                                                                                                             ABA98547;
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             (BIOW-) BIOWINDOW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human zinc finger protein 9.79 polypeptide, diseases e.g. cancer and HIV infection.
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                                                                                                                                                                                                                                                                                                                                         Similarity
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CR; primer; ss.
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llarity 78.3%;
Conservative
                                                                                                                                                                   growth factor binding protein 16.17 PCR primer #1.
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             GENE DEV INC
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Pred. No. 2.3e
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Best Local (
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                                      The invention relates to a novel polypeptide, argininoacyl tRNA synthetase 12.87, and the polypucleotide encoding it. The polypeptide is useful for treating various diseases, such as malignant tumours, inflammations, immunological diseases, haemopathy and human immunodeficiency virus (HIV) infection. The invention also discloses the antagonist resisting the polypeptide and its treatment effect, and application of the polymucleotide. The present sequence represents a PCR primer used to amplify the argininoacyl tRNA synthetase 12.87 gene of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Insulin-like growth factor binding protein 16.17 and encoding
polynucleotide, used in diagnosis and treatment of cancer.
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                                                                                                                                                                                                        New polypeptide argininoacyl tRNA synthetase 12.87 and encoding polynucleotides, useful for treating malignant tumors, inflammations, immunological diseases, hemopathy and human immunodefiency virus
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                              invention
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lmmunological disease; haemopathy; human immunodeficiency virus; HIV;
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AAZ95163
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                               Query Match
Best Local Similarity
Matches 18; Conserv
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Best Local Similarity
Matches 18; Conserv
                                                                                                                           molecules, which comprise human UGT2B sequence polymorphisms (see AR295051-295110). Probes which detect the UGT2B locus polymorphisms can be used to detect altered UGT2B metabolism of a substrate in an individual. The nucleic acid molecules comprising a human UGT2B sequence polymorphism can be used in screening assays for genotyping individuals, also to predict their rate of metabolism of UGT2B substrate, potential drug-drug interactions and adverse side effects. The polymorphisms can be used as single nucleotide polymorphisms (SNPs) for detecting genetic linkage related to phenotypic variation in activity or expression of UGT2B protein. The polymorphism containing nucleic acid molecules may also be used for generating genetically modified non-human animals and for obtaining site specific gene modification in cell lines
                                                                                                                                                                                                                                                                                                                                 This sequence represents a primer used to sequence polymorphic fragments of the human UDP-glucuronosyltransferase 287 (UGT287) gene. UDP-glucuronosyltransferase 287 (UGT287) gene. UDP-glucuronosyltransferases (UGT8) are a family of enzymes that catalyse the glucuronic acid conjugation of a wide range of endogenous and exogenous substrates. The UGT28 gene subfamily encode steroid metabolizing isoforms in the liver. Alteration of the expression or function of UGTs may effect drug metabolism. The invention relates to non-chromosomal nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel human UDP-glucuronosyltransferase sequence, polymorphisms for genotyping individuals to predict rate of metabolism of substrates for identifying potential drug interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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drug interaction; detect; human; single nucleotide polymorphism; primer;
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                                                                                                 Sequence
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4023 AAAGAGAGAAAACAAAATGTTAT 4045
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                                 Conservative
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                                                                                                 A; 2 C; 1 G; 5 T; 0 U; 0 Other;
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78.3%;
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Pred. No. 2.3e
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                             Score 15; DB Pred. No. 2.3e 0; Mismatches
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ABK99281
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AAD46030
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                                                                                                                                                                                                                                                           Query Match
Best Local
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Synthetic
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                                                                                                                                                                                                                                                                                                                        glucuronidated drug that may be used in treating cancer patients. It is also useful in determining persons at risk for epirubicin toxicity, in reducing or eliminating side effects associated with epirubicin treatment, and in ways of increasing the efficacy of dosage regimens. The present sequence is a primer used for sequencing human UGT2B7 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining the accommended determining
                         Hepatitis C virus; HCV; NS5B replicase; ss; RNA polymerase
                                                                                                                             ABK99281 standard; RNA;
                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 16 A; 2 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an UDP-glucoronosyl transferase (UGP) enzyme, UG72B7. The invention also relates to compositions and methods for optimising UG72B7 substrate dosings and for predicting UG72B7 substrate toxicity. The method is useful in determining the dose of a UG72B7-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  patient.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; UDP-glucoronosyl transferase; UGT; UGT2B7; toxicity; cancer;
                                                   Hepatitis C virus (HCV) NS5B replicase RNA synthesis template
                                                                            21-OCT-2002
                                                                                                       <u> АВК99281;</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 53; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-691534/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy; epirubicin; cytostatic; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human UGT2B7
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                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the dose of a UGT2B7-glucuronidated drug fetermining the level of UGT2B7 activity or
                                                                                                                                                                                                                                               Conservative
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Pred. No. 2
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                                                                                                                                                                                                                                                           DB 1;
2.3e+03;
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                                                                                                                                                                                                                                                                      Length 24;
                                                                                                                                                                                                                                                 Indels
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RESULT 2901
ABS57119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer is two or three nucleotides, and the template and primer do not form a stable duplex in solution in the absence of the MCV NSSB protein. The complex is useful for detecting HCV replicase activity and permits establishment of sensitive RNA-dependent RNA polymerase assays to screen and evaluate antiviral inhibitors and to improve the specificity and cefficacy of the inhibitors. The complex is also useful in the development of a reliable system for determining kinetic and thermodynamic constants of HCV NS5B-catalysed nucleotide incorporation and investigation of mechanistic inhibitors for mis-incorporation or chain termination. Specifically, the short RNA template and primer pairs are useful in Screening assays which are used for determining kinetic, thermodynamic and mechanistic properties of NS5B replication and ultimately in the development of inhibitors of NS5B newly identified inhibitors of replicase activity may be used for developing anti-HCV pharmaceuticals. Sequences ABK99271-ABK99296 represent HCV NS5B replicase RNA synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a replicase complex comprising a hepatitis C virus (HCV) NS5B replicase protein, a linear nucleic acid template and complementary nucleic acid primer which is annealed to the 3' terminus the template, where the template is at least three nucleotides and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel replicase complex comprising hepatitis C virus NS5B replicase, a 3 nucleotide-long template to which a 2 nucleotide-long primer is annealed, and template and primer which do not form a stable duplex in the absence
CN1352095-A
                                                                Human; ss; shear protein 8.91; tumour; haemopathy; HIV; PCR; primer; human immunodeficiency virus infection; immunological disease; inflammation; RT-PCR; reverse transcriptase PCR.
                                                                                                                                                                        30-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 6; 17pp; English.
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                                Homo sapiens
                                                                                                                                                                                                                                            ABS57119 standard; DNA; 24 BP
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                                                                                                                                  Huma shear protein 8.91 RT-PCR
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                                                                                                                                                                                                                                                                                                                                                                 5774 GCCGGCCTGCCTGCCTGCC 5796
                                                                                                                                                                                                                                                                                                                                                                                                   18;
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Similarity 78.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15; DB 1;
Pred. No. 2.3e+03;
                                                                                                                                    primer
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ID AAX84260 standard; DNA; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein 8.91, polymucleotides encoding the polypeptide and producing the polypeptide by recombinant DNA technology. The present invention also discloses applying the polypeptide in treating various diseases, such as malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present invention also discloses the antagonist resisting the polypeptide and its treatment effect. The present invention also discloses application of the polymucleotides encoding human shear protein 8.91. The present sequence is a reverse transcriptase (RT)-PCR primer used to isolate nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New human shear protein 8.91 polypeptide for treating malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention discloses a new kind of polypeptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-644454/70
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                                                           (KYOW )
                                                                                             15-DEC-1997;
                                                                                                                     14-DEC-1998;
                                                                                                                                             24-JUN-1999.
                                                                                                                                                                       WO9931239-A1
                                                                                                                                                                                               Homo
                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                Nck associated protein 1; Napl; human; apoptosis; Alzheimer's disease;
therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                    PCR primer for human Nck associated protein 1 coding sequence
                                                                                                                                                                                                                                                                                              08-SEP-1999
                                                                                                                                                                                                                                                                                                                     AAX84260;
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                                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
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                                                          KYOWA HAKKO KOGYO KK
SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BODE GENE
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L8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human shear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 16 (disclosure); 32pp; Chinese.
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WPI; 1999-395181/33

Sakaki Y;

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RESULT 2903
ABZ2353
XX ABZ2353
XX ABZ2353
XX ABZ2353
XX PROPR-
XX Replica
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Best Local S
Matches 18
The specification describes a method for determining whether a product a gene is involved in preventing a replication error in a cell. The method comprises providing the cell with a specific inhibitor for the product and determining the level of functional expression of a marker gene in the cell, where the level of expression of the marker gene is dependent on the occurrence of the replication error. The method is use for determining whether a product of a gene is involved in preventing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents a PCR primer used to isolate DN human Nck associated protein 1 (Wap1) of the invention. apoptosis. The protein can be used in the investigation, treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                                                                                                                                                                           Determining whether a product of a gene is involved in preventing a replication error in a cell comprises providing a specific inhibitor the product and determining the level of expression of a marker gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Replication
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                                                                                                                                                                                   Example 1; Fig 3; 47pp;
                                                                                                                                                                                                                                                                                                                                                                           Plasterk
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TIJSTERMAN M.
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                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          development; somatic instability;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    base represents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .4e+03;
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Query Match
Best Local S
Matches 18
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Best Loc
Matches
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                                       Sequence
                                                                    This invention relates to the goat beta-casein gene, represented by AAF77439. There are 18 positions in the 5'-flanking region, 21 positions in intron sequences and 1 position in exon VII that conflict with the sequence reported for this gene by Roberts et al (1992). The present sequence represents a fragment of the casein gene in which a conflict occurs when compared to the Roberts et al (1992) sequence for the same
                                                                                                                                           Disclosure; Page 34; 41pp; Chinese
                                                                                                                                                                DNA molecules containing goat beta-casein
                                                                                                                                                                                                                                                                                                                                                                           Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF77454 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   replication error in a cell. The identified genes are useful for developing diagnostic tools, or as targets for drug development to manipulate cells on the basis of the presence or absence of function the gene. ABZ23535-36 represents fragments of plasmids used to detect
                                                                                                                                                                                                         Huang
                                                                                                                                                                                                                                                 28-MAY-1997;
                                                                                                                                                                                                                                                                      28-MAY-1997;
                                                                                                                                                                                                                                                                                          01-DEC-2000
                                                                                                                                                                                                                                                                                                               TW413702-A
                                                                                                                                                                                                                                                                                                                                                                 conflict
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Intron
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          somatic instability, in the
                                                                                                                                                                                                                              (NASC-)
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                                                                                                                                                                                                                                                                                                                                                                                                                   beta-casein;
                                                                                                                                                                                                                                                                                                                                                                                               hircus
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18; Conserv
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          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                      fragment 4 of
                                        BP; 0
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                                                                                                                                                                                                                                                 97TW-00107261.
                                                                                                                                                                                                                                                                      97TW-00107261.
                                                                                                                                                                                                                                                                                                                              /note= "Conflict occurs at this location when beta-casein gene given by Roberts et al (1992)
                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers replace(12. .14,TC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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72.0%;
                                        4 C;
        78.
        .3%;
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Score 15; DB pred. No. 2.4e 0; Mismatches
                                        G; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                       beta-casein
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                                       u; o
         DB 1;
2.4e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4e+03
                                        Other;
                    Length
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RESULT 2906
AAS20672/c
ID AAS2067
XX AAS2067
XC AAS2067
XC O9-APR-
XX
DE Human z
XX
KW Cytokin
                                                                                                                                                                                                                                                                                                                  RESULT 2905
AAI73048/c
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                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                        This sequence represents a molecular scaffold which may be used in a composition of matter comprising multiple fluorophores. The fluorophores are bound to the molecular scaffold at separate predetermined positions, to permit fluorescence energy transfer between two fluorophores. The fluorophores are characterized by maximum emission wavelength of one being greater than the minimum excitation wavelength of the other. The composition is useful for determining whether a preselected nucleotide residue is present at a predetermined position within a nucleic acid. It is also useful in multicomponent analysis including multiplex biological analysis, and identifying multiple single nucleotide polymorphisms. The presence of a number of given nucleotide residues is determined
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Composition of matter useful for multi-component analyses, comprises multiple fluorophores bound to molecular scaffold at preset positions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-SEP-2000;
31-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-SEP-2001; 2001WO-US028967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200222883-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Molecular scaffold; fluorophore; fluorescence; energy transfer; emission wavelength; excitation wavelength; multiple; single nucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scaffold oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI73048 standard; DNA;
 Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor;
                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-575158/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ju J,
                                                             09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYCO ) UNIV
                              Human zalphall Ligand sequencing primer 2C7764b
                                                                                                                       AAS20672
                                                                                                                                                                                                                                                                                                                                              simultaneously by the composition of the invention
                                                                                                                                                                                                                           4017
                                                                                                                                                                                              26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fluorescence energy transfer between two fluorophores.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Li Z,
                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                           GAGAAAAAAGAGAGAAAACAAAA 4039
                                                                                                                                                                                                                                                                                                               26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                       standard;
                                                                                                                                                                                              GAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 43; 113pp; English.
                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-00658077.
2001US-0309156P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COLUMBIA NEW YORK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tong A,
                                                                                                                     DNA;
                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Russo JJ;
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Pred. No. 2.4e.
0; Mismatches
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                                                                                                                                                                                                                                                                                      DB 1;
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RESULT 2907 ABX93461/c

ABX93461 standard; DNA; 26

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26

GAAAAAAAAAAAAAAAAAAAA

4017 GAGAAAAAGAGAGAAAACAAAA 4039

Query Match Best Local S Matches 18

Similarity

78.3%;

Score 15; DB 1; Pred. No. 2.4e+03;

Length 26;

Conservative

<u>..</u>

Mismatches

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Indels

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0

SXXXXXXXXXXXXX

27-MAY-2003 (1 LS147-specific

polynucleotide sequencing

related universal primer #1

(first entry)

LS147; cancer; lung cancer; gene therapy; cytostatic;
primer; EST clone; expressed sequence tag clone.

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The present invention relates to the isolation of a novel cytokine, CC zalphall Ligand and the polynucleotide encoding it. The invention also CC gives the sequence for the zalphall receptor and the polynucleotide encoding it. The zalphall rigand polypeptide stimulates proliferation of CC entural killer (NK) cells or NK cell progenitors, the activation of NK CC cells, proliferation of T-cells, proliferation of B-cells stimulates an antibodies response in a mammal, and CC reduces proliferation of B-cells stimulates an antibodies. The zalphall Ligand polypeptide is also useful in preparing antibodies that CC bind to zalphall Ligand epitopes. The zalphall Ligand golypeptide is also useful in preparing antibodies that CC inhibitors of its activity, to enhance the generation of anti-tumour CC inhibitors of its activity, to enhance the generation of anti-tumour CC responses with or without the infusion of donor lymphocytes, and to CC activate or stimulate the immune system. The present sequence represents CC human zalphall Ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-MAR-1999;
11-MAR-1999;
01-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    natural killer cell proliferation; T-cell proliferation; B-cell proliferation; anti-tumour response; immune syste immunostimulant; cytostatic; human; sequencing primer; s
                                                                                                                                                                                                                                                                                                                                                                                                 New zalphall ligand polypeptides and polynucleotides, useful for stimulating proliferation, activation, differentiation and/or induction of inhibition of specialized cell function, or for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-040208/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novak JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-OCT-2001.
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Sequence 26
                                                                                                                                                                                                                                                                                                                                                  Example 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                  Col 139; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Johnston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Presnell
 BP; 0 A; 1 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-00522217.
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99US-0123904P.
99US-0142013P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sprecher CA,
Nelson AJ, I
25 T; 0 U; 0 Other;
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ABX12469/c
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Best Local
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                                                                                                                              Coxsackie virus strain VD2921; diabetogenic coxsackie B virus-4; strain VD2921; VP1; VP3; VP4; P2A; P2B; P2C; P3A; P3B; P3C; diabetes; diabetogenic enterovirus; beta cell loss; blindness; renal failure; leg amputation; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a purified polypeptide or its fragment derived from the LS147 gene capable of selectively hybridising to the nucleic acid of the gene and has at least 50% identity with the polymucleotide. The LS147 polypeptide is useful for preparing a composition for treating cancer, e.g. lung cancer using gene therapy. This sequence represents a universal primer used to sequence LS147 expressed sequence tag (EST)-
                          20-JUN-2001; 2001SE-00002198.
                                                19-JUN-2002;
                                                                                                            Coxsackievirus.
                                                                                                                                                                                   Coxsackie B
                                                                                                                                                                                                      10-MAY-2003
                                                                                                                                                                                                                          ABX12469;
                                                                                                                                                                                                                                              ABX12469 standard;
                                                                                                                                                                                                                                                                                                                                                                                Sequence 26
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Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUN-1998;
       (INNO-) INNOVENTUS PROJECT AB
                                                                     27-DEC-2002
                                                                                       WO2002103060-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New LS147 polypeptide, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-341045/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RUSS/)
(STRO/)
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                                                                                                                                                                                                                                                                                                                    4017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lung
                                                                                                                                                                                                                                                                                                26
                                                                                                                                                                                                                                                                                                                                         18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COHEN M.
COLPITTS T L.
FRIEDMAN P N.
KLASS M R.
RUSSELL J C.
STROUPE S.
                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Page 39; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BILLINGEL
                                                                                                                                                                                                                                                                                                                   GAGAAAAAAGAGAGAAAACAAAA 4039
                                                                                                                                                                                                                                                                                                GAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer.
                                                                                                                                                                               virus 4 (CBV-4) strain VD2921, PCR primer dT26V
                                                                                                                                                                                                                                                                                                                                                                                BP; 0
                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cohen M,
Stroupe S;
                                                2002WO-IB003278
                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-0048810P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-00092296.
                                                                                                                                                                                                                                                                                                                                                                                P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ъ.
                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                0.2%;
                                                                                                                                                                                                                                                                                                                                                                                1 C; 0
                                                                                                                                                                                                                                              27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Colpitts TL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                      <u>..</u>
                                                                                                                                                                                                                                                                                                                                                                               G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                     Score 15; DB 1; Le
Pred. No. 2.4e+03;
0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                preparing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a composition
                                                                                                                                                                                                                                                                                                                                                           Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PN,
                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Klass MR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating
                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                 CBV-4;
P3D;
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                                                                                                                                                                                                                                                      ABN83378
                                                                                                                                                                                                                                                                                                          Best Loc
Matches
                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                      83
                                                                                                                                                        Key
                                                                                                                                                                                                                                                                                                                                    Sequence
                         WPI; 2002-437469/47
                                       Dietmaier W;
                                                                                   13-NOV-2001; 2001EP-00126930
                                                                                                               EP1207210-A1
                                                                                                                                               modified_base
                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                    15-AUG-2002
                                                                                                                                                                                                                                 ABN83378;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tuvemo
                                                   (HOFF ) ROCHE DIAGNOSTICS GMBH (HOFF ) HOFFMANN LA ROCHE & CO
                                                                       15-NOV-2000; 2000EP-00124897.
                                                                                                  22-MAY-2002.
                                                                                                                                                                                        Mononucleotide repeat locus; human; BAT25; probe; microsatellite; tumour;
                                                                                                                                                                                                     Mononucleotide repeat locus BAT25 probe #1
                                                                                                                                                                                                                                              ABN83378 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymerase chain the diabetogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                         4011 TAAAATGAGAAAAAAAGAGAGAAAAACAA 4037
                                                                                                                                                                                                                                                                                27
                                                                                                                                                                                                                                                                                                          19;
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                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                BAAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 44; 79pp;
                                                                                                                                                                                                                                                                                                                                    B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Frisk GE,
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                    0 A;
                                                                                                                                                Location/Qualifiers
                                                                                                                             /note= "Labelled with Fluorescein"
                                                                                                                                  /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                            reaction and primers for detecting nucleic coxsackie B virus-4 strain VD2921.
                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                    C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yin
                                                                                                                                                                                                                                              29
                                                                                                                                                                                                                                                                                                                                                                                                                                 polymerase chain reaction (PCR) and primers for
                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
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                                                                                                                                                                                                                                                                                                         Score 15; DB
Pred. No. 2.5e
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                    26 T; 0 U; 1 Other;
                                                    Ŧ
                                                                                                                                                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                                 5e+03
                                                                                                                                                                                                                                                                                                                      Length 27;
                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          B virus-4 (CBV-4)
                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acids
                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                    P3B,
                                                                                                                                                                                                                                                                                                          0
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Analyzing repeat sequences in DNA using a probe which hybridizes to adjacent repetitive and non-repetitive regions and determining hybrid melting point is useful to detect microsatellite instability such as

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RESULT 2910
AAS63441
ID AAS6344
XX AAS6344
XX Oligonu
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Best Local Simi
Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                               28-MAR-2000; 2000US-0192699P.
26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-02013906P.
08-DEC-2000; 2000US-0213906P.
11-DEC-2000; 2000US-0254392P.
11-DEC-2000; 2000US-0256395P.
12-JAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-00820279.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method for analysing a target nucleic acid consisting of repetitive and non-repetitive sequences. The method comprises hybridising a polynucleotide probe comprising a segment complementary to an adjacent repetitive region, where the second segment consists of a defined number of repeats, and determining the melting point temperature of the hybrid. The method is used to analyse microsatellites, especially microsatellite instability, particularly as a means for detecting hereditary tumours. Alternatively, the method is used to identify an individual in a population. The present sequence is a probe for hypertical repeat locus BAT25, and was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide-nanoparticle probe; dia nucleic acid detection; nanostructure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hereditary cancer
                                                                                                     Detecting and separating nucleic acid, comprises reaction with nanoparticles to complementary to parts of the target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide-nanoparticle probe #63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS63441 standard; DNA; 30 BP
                                                      Example 24; Fig 44; 404pp;
                                                                                                                                                                                                                   WPI; 2001-656926/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAR-2001; 2001WO-US010071.
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                                                                                                                                                                                                                                                                                                                                                   (NANO-) NANOSPHERE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAAAAAAAAAAAAAAAT
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                                                                                                                                                                                                                                                                    Park S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 26 A; 2 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                       Letsinger |
Park S, Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7; 19pp;
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                                                                                                                                                                                                                                                                    RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
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                                                      English.
                                                                                                                                                                                                                                                                                               Mucic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27
                                                                                                                                                                                                                                                                                               RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ŏ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnostic;
re; biochip;
                                                                                                                              , useful e.g. for diagnosis, that carry oligonucleotides
                                                                                                                                                                                                                                                                                               Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ģ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    forensic analysis;
biofilter; drug delivery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                               Elghanian
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The invention relates

to a method for detection of

nucleic

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RESULT 2911
AAS10385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC having at least 2 portions, comprising treatment with nanoparticles that C carry oligonuclectides complementary to at least 2 parts of (1), where CC detectable change caused by hybridisation of the oligonuclectide to (1) of sobserved. The method is used to detect (or to separate) specific (1), CC e.g. for diagnosing a wide variety of diseases, sequencing, in forensic can large to the carrivatised nanoparticles are also useful for preparing consostructures useful, for example, as blochips, biofilters, mechanical conjugates, separation membranes, chemical sensors, in computers, and for CC drug delivery. Very stable nanoparticle-oligonuclectide conjugates can be produced, allowing their direct use (as probes) in polymerase chain creation, i.e. they survive multiple heating/cooling cycles so do not consider, without the need for special equipment, making possible rapid coligonucleotide-nanoparticle probes, and related sequences, used in the cc method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 18
Detecting a nucleic acid useful in e.g. diagnosing genetic, bacterial or viral diseases, by contacting the nucleic acid with oligonucleotides attached to nanoparticles and having sequences complementary a portion of the nucleic acid.
                                                                                                                                                             13-JAN-2000; 2000US-0176409P
26-APR-2000; 2000US-0200161P
26-JUN-2000; 2000US-00603830
12-JAN-2001; 2001US-00760500
                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                     Nanoparticle; cyclic disulphide-oligonucleotide; DNA detection; DNA isolation; genetic disease; bacterial disease; viral disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide-cyclic disulphide linker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS10385 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 30
                                                                                                          Mirkin
                                                                                                                                                                                                                                                                                                                                                   misc_feature
                                                                                                                                                                                                                                                                                                                                                                     Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-OCT-2001
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                                                                                                                                                                                                                                                            19-JUL-2001.
                                                                                                                                                                                                                                                                                     WO200151665-A2
                                                                                                                                                                                                                                                                                                                                                                                                                       forensic science;
                                                                                                                                      (NANO-) NANOSPHERE INC.
                                                                                                                                                                                                                                  12-JAN-2001; 2001WO-US001190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3278
                                                                    2001-451868/48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 23
                                                                                                          Letsinger RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                              27
                                                                                                                                                                                                                                                                                                                          /*tag= a
/note= "A
                                                                                                                                                                                                                                                                                                               moiety"
                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                              is covalently
                                                                                                           Mucic
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                                                                                                                                                                                                                                                                                                                                                                                                                       testing; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                           RC,
                                                                                                              Storhoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                             linked
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                                                                                                                                                                                                                                                                                                                             cyclic-disulphide
                                                                                                            Elghanian
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RESULT 2912
ABK65048
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence represents a cyclic disulphide linked oligonuclectide which CCC may be coupled with colloidal gold particles (nanoparticles) and used to CC demonstrate the method of the invention. The invention relates to CC demonstrate the method of the invention. The invention relates to CC isolating or detecting a nucleic acid of interest, in a mixture of CC nucleic acids, by binding it to 2 or more complimentary nucleotides which CC colloidal gold) are used to both isolate and detect (e.g. by linking the particle to a fluorescent probe) the resultant complex. The methods are CC useful for detecting nucleic acids, natural or synthetic, and modified or CC unmodified. The methods may also be applied in the diagnost of genetic, bacterial and viral diseases, in forensics, in DNA sequencing, for CC paternity testing, for cell line authentication, and for monitoring gene CC therapy. The methods are further useful in research and analytical CC specific pathogens, for quick identification of an infection to assist in CC drug prescription, and in homes and health centres for inexpensive first-CC line screening. The methods, which are based on observing colour change CC with the naked eye, are cheap, fast, simple, robust (reagents are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 18
                                                                                                                                                                              11-AUG-2000;
08-DEC-2000;
11-DEC-2000;
12-JAN-2001;
28-MAR-2001;
           Detecting nucleic acid, useful for diagnosis of genetic, viral or bacterial disease, comprises hybridizing nanoparticles with attacoligonucleotides to nucleic acid and detecting change brought about
                                                                                                                                                                                                                                                                                                          07-MAR-2002.
                                                                                                                                                                                                                                                                                                                                       WO200218643-A2
                                                                                                                                                                                                                                                                                                                                                                                                                  Nanoparticle-oligonucleotide; nanofabrication;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Nanoparticle-oligonucleotide #68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK65048 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     line screening. The methods, which are with the naked eye, are cheap, fast, si stable), do not require specialised or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 24; Fig 44; 323pp; English.
                                                                                                                                                                                                                                                                            10-AUG-2001; 2001WO-US025237.
                                                                        WPI; 2002-258024/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            instrumentation is required
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                   NANOSPHERE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAAAAAAAAAAAAAAGCAGACC 27
                                                                                                     Letsinger RL,
Garimella V,
                                                                                                                                                                                               ; 2000US-0224631P.
; 2000US-0254392P.
; 2000US-0255235P.
; 2001US-00760500.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.2%;
ilarity 78.3%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B₽;
                                                                                                                                                                                2001US-00820279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                     Mucic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15; DB 1; Le
Pred. No. 2.7e+03;
D; Mismatches 5;
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                                                                                                      Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3300
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                                                                                                      s
                                                                                                                   Storhoff JJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expensive equipment,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
            articles with attached change brought about by
                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                   Elghanian
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CC The invention relates to a method of detecting a nucleic acid (NA) having CC at least 2 portions comprising: (a) providing nanoparticles (NP) with CC attached oligonuclectides (CCN), where CON has a sequence complementary CC to the sequence of NA; (b) contacting NA and NP under conditions CC effective to allow hybridisation of CGN with NA; and (c) observing a CC detectable change brought about by hybridisation of CGN with NA. The CC method is useful for detecting a nucleic acid, separating a selected CC nucleic acid from others and methods of nanofabrication. Detecting CC analytes such as nucleic acids and proteins are useful for the diagnosis CC fgenetic, bacterial and viral diseases. The CGN NP conjugates that use CC cyclic disulphide linkers improve the sensitivity of diagnostic assays. In particular assays using CGN-NP conjugates prepared using linkers (comprising a steroid residue attached to a cyclic disulphide have been CC found to be approximately 10 times more sensitive than assays employing CC conjugates prepared using alkanethiols or acyclic disulphides as the CC linker. The CGN-NP conjugates are stable allowing them to be used CC directly in PCR solutions. Therefore conjugates added as probes to a DNA CT of the conjugates of the PCR and are still able to detect the amplicons without opening the tubes and causing contamination. ABK64981-ABK65055 represent nanoparticle-oligonucleotides of the invention
DRX PRARA PRARA PRAKA PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 2913
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Best Local S
                                                                                                                                                                                                                                 08-DEC-2000;
08-DEC-2000;
11-DEC-2000;
11-DEC-2000;
12-JAN-2001;
28-MAR-2001;
09-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid detection method; nanoparticle-oligonucleotide conjugate; nanoparticle; viral RNA detection; bacterial DNA detection; fungal DNA detection; nanoprobe conjugate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-DEC-2001; 2001WO-US046418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS64686;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS64686 standard;
                                                                                                                                                                                                        10-AUG-2001;
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                                        1 CA,
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                                                                                                                                              NANOSPHERE INC.
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                                                       Letsinger RL, N
Garimella V, Li
                                                                                                                                                                                                  2000US-0254392P.
2000US-0254318P.
2000US-0255236P.
2000US-0255236P.
2001US-00760500.
2001US-00820279.
2001US-0282640P.
2001US-00927777.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 23 A; 4 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detection method associated polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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78.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                    Mucic RC, S
i Z, Park S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27
                                                                                   Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 30;
                                                                                   Elghanian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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WPI; 2002-608256/65

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RESULT 2914
AAL61658
AID AAL6165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes a method of detecting (MI) a nucleic acid having two portions, involving providing nanoparticles having oligonucleotides attached to it, which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and nanoparticles, to allow hybridisation of oligonucleotides with two or more portions of nucleic acid, and observing a detectable change brought about by hybridisation. (MI), nanoparticles (I), nanoparticle-oligonucleotide conjugates (II) and the aggregate probe are useful for detecting two or more nucleic acids (from a biological source) having at least two portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification. (II) is useful for preparing a nucleic acid bound to an electrode surface. (I) and (II) are useful for fabrication, and for separating a selected nucleic acid having two portions from other nucleic acids. (I), (II) and the aggregate probe are useful for detecting an analyte (especially polymucleotide used to demonstrate the method of the invention
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide #19 used in the nucleic acid detection method
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                                                                                                                                                                                                                                                                                                                                                                                                                             09-OCT-2001; 2001US-0327864P.
07-DEC-2001; 2001US-00008978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid detection; fabrication; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAL61658
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                                                                                                       nanoparticles to allow hybridization,
                                                                                                                                     Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and
                                                                                                                                                                                                                                      WPI; 2003-430409/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-OCT-2002; 2002WO-US032088.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
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                                                                                                                                                                                                                                                                                                  Taton TA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fig 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     442pp; English.
                                                                                                                                                                                                                                                                                                     Mirkin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.2%;
78.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15; DB 1;
Pred. No. 2.7e+03;
                                                                                                          and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                       observing
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                                                                                                       detectable change
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Example 24; Fig 44; 467pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  portions. The method involves providing nanoparticles having colligonuclectides attached to it which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and comparticles to allow hybridisation of oligonucleotides with two or more partions of nucleic acid and observing a detectable change brought about by hybridisation. The method and aggregate probes are useful for detecting two or more nucleic acids (from a biological source) having at least two portions such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic or structurally modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification. The invention is useful for preparing a nanoprobe conjugate for detecting an analyte and for detecting a selected nucleic acid having two portions from other nucleic separating a selected nucleic acid having two portions from other nucleic acids. The present sequence is an oligo used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                            misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                      Human leukocyte adhesion protein; p150,95 alpha subunit gene; herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 23 A; 4 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligomer LAP312 for forming triplex with HUMINT02 target duplex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ30395 standard; DNA;
                    modified_base
                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ30395,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of detecting a nucleic acid having two
                                                                            modified_base
                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                            misc_feature
                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3278 AAGAAGAAAAATGAAACCAGACC 3300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                       13
                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                         /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                               /note= "OTHER= N6 methyl-8-oxo
/mod_base=
                                       note=
                                                   mod.
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                                                                                                                                   10. .18
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                                     _base= OTHER=
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                                                                                                      inverted_polarity_region
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Pred. No.
                                                                                            comments"
                                        8
                                                                                                                                               N6 methyl-8-oxo 2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                     dimer synthon linkage"
                                        methyl-8-oxo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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                                         ń
                                                                                                                                                                                                                                            2' deoxyadenine"
                                                                                                                                                                                                                                                                                                 2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 2916
AAQ38707
                                                                                                                                                                                                                                         The synthetic oligomer is capable of forming a triplex at physiological CC pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human leukocyte adhesion protein p150, 95 alpha subunit gene (HMMNT02) CC beginning at nucleotide 677 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in CC diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV; HER; HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays CC may be carried out without subjecting the test specimen to harsh CC conditions. The oligomer contains an inverted polarity region formed from an o-xyloso dimer synthon. The linking gp. is o-xyloso (nucleotides are coupled through a xylene residue to form the dimer synthon. This additional modifications may render the oligomer stable to nuclease activity. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448.
                                                                                                                                                              Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-NOV-1990;
18-JAN-1991;
08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
 25-MAR-2003
                         AAQ38707;
                                                 AAQ38707
                                                                                                                                                                                                                      Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 12; Page 70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JUN-1992.
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                                                                                                                                         6682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      malignancy and inflammation.
                                                                                                                 1 TAATTTTAATTATATAT 18
                                                                                                                                 TTATTTTTATTTATATAT 6699
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                                                   standard;
                                                                                                                                                                                                                     BP;
                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Krawczyk S,
(revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90US-00617907.
91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
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17
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                                                   RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77pp; English.
                                                                                                                                                                              0.2%;
                                                                                                                                                                                                                     0 C; 0 G; 11 T; 0 U;
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e= "OTHER= N6
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                                                                                                                                                                               Score 14.8;
Pred. No. 1.
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N
                                                                                                                                                                   Mismatches
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                                                                                                                                                                               1.8e+03
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                                                                                                                                                                                          DB 1; Length 18;
                                                                                                                                                                                                                     0 Other;
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AAQ92177
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Best Local S
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                                     10-NOV-1994;
                                                                                                                Synthetic.
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            12-NOV-1993;
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                                                                                      WO9513397-A1
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This sequence was used as a PCR primer in order to add a polyA tail to the 3'end of the highest specific activity selected oligonucleotide in order to form a first strand. The primer is comprised of a 5'known sequence and a 3' polynucleotide portion corresp. to the polynucleotide tail of the first strand. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide binding; nucleotide binding; DNA detection; binding DNA treatment; diagnosis; testing; assay; Candida; papillomavirus; cytomegalovirus; pstein-Barr ir ir rhinovirus; hepatitis virus; liver disease; human immunodeficiency virus; herpes simplex virus; human immunodeficiency virus; HIV; ALDS; influenza virus; genetic disease; genetic abnormalities.
                                                                                                                                                                                                                                                                          Primer; polymerase chain reaction; amplify; mutant; K-ras; flanking region; amplification; probe; detection; sputum; obenign; malignant; neoplasm; lung; lung cancer; head; neck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determn. of oligo:nucleotide(s) with specific activity for a - for use in therapeutics, diagnostics and research reagents.
                                                                                                                                                                                                                                                                                                                                                                                         p53 detection probe,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 1 A; 0 C; 3 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 27; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ92177 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1993-101001/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JUL-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4460 GGACTTTTTTTTTTTTT 4477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric primer for adding poly A tails
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
9308-00152313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91US-00755485
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                                                       94WO-US012947
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                                                                                                                                                                                                                                                                                                                                                                                               273 CGT to TGT)
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                             sputum; diagnosis;
                                                                                                                                                                                                                                                                                neck; ss
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RESULT 2918
AAQ83415
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequences given in AAQ92112-211 are probes which were used in the detection of a mutant p53 gene sequence. The DNA to be detected is amplified using PCR and then these probes which are pref. labeled using 32-P gamma-ArP are used to detect the mutant sequences. The primers and probes given in AAQ92098-219 are used in the method of the invention for detecting mammalian target DNA in sputum samples. Analysis of the target DNA is used to diagnose benign or malignant neoplasms of the lung. It is also useful for screening people at high risk or for monitoring progress of treatment of lung neoplasms. The method is based on the discovery that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting target nucleic acid in mammalian sputum diagnosis of lung neoplasia involving mutation(s) or p53 tumour suppressor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mutant target DNA associated with lung cancer is present at detectable levels in sputum. Cells shed into sputum from head and neck cancers may also be detected
Antisense nucleic acid hybridising with an comprising the genes c-jun, jun-B or c-fos,
                                                             Use of antisense c-jun, c-
treating neuronal injury,
                                                                                                                                                                                                                                                                                                                           c-jun,
                                                                                                                                                                                                                                                                                                                                                    c-fos antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
20-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ83415 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 34; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-194114/25.
                                      Claim 2;
                                                                                                       WPI; 1995-066896/09.
                                                                                                                              Schlingensiepen G,
                                                                                                                                                                                  10-JUL-1993;
                                                                                                                                                                                                          06-JUL-1994;
                                                                                                                                                                                                                                    19-JAN-1995.
                                                                                                                                                                                                                                                             WO9502051-A2
                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                               phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            μ
                                                                                                                                                                                                                                                                                                                         c-fos; jun-B; neuronal injury; cell death; neoplasm; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                          BIOGNOSTIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTGAGGTGTGTGTTTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTGAGATTTGTGTTTTGTG 7326
                                     Page 61; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                  93EP-00111059
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                                                                                                                                                                                                                                                                                                                88
                                                                                                                                                          GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.9%;
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                                                                                                                              Schlingensiepen R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
                                                               c-fos or jun-B
y, degeneration,
                                                                                                                                                                                                                                                                                                                                                                                                                                             18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 U;
                                                                nucleic acids - for , cell death and/or r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8e+03;
                                                                                                                            Schlingensiepen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
area of the mRNA and/or DNA, expression of which plays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 18;

    particularly for 
in the K-ras oncogene

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                               ζ,
                                                                neoplasms
                                                                            preventing
                                                                                                                               Brysch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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4477 0

Matches Query Match Best Local S

16;

Conservative

Similarity

0.2%;

Pred.

1.8e+03

Mismatches ö.

Score 14.8; DB 1;

Length 18; Indels

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Gaps

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  causal role in neuronal injury, degeneration, cell death and/ or neoplasms, can be used to prevent and treat such conditions. c-jun antisense sequences are described in AAQ83267-321 and AAQ83440-43; jun-B antisense sequences are described in AAQ83322-63 and AAQ83444-45; and c-for antisense sequences are described in AAQ83364-439 and AAQ8346-51. Proferably the antisense sequences are phosphorothicate oligonucleotides since these are not destroyed as fast by endogenous factors as naturally occurring molecules. (Updated on 25-MAR-2003 to correct PN field.)
                                        determining an oligonucleotide having specific activity for a target biomolecule. The method comprises assaying a set of randomised oligonucleotides for activity against a target biomolecule, separating active from inactive oligonucleotides and recovering, amplifying and determining the nucleic acid sequence of the active oligonucleotides. oligonucleotides can be used for therapeutic, diagnostic and research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Determination; oligonucleotide; specific activity; therapy; target biomolecule; randomised oligonucleotide; diagnosis; chimeric; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            First
                                                                                                                                                                                       Determination of oligo-nucleotide with specific activity for target biomolecule - using set of randomised oligo-nucleotide(s).
                                                                                                                                                                                                                                                                                                                         05-SEP-1991;
04-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT96107 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 7 C; 0 G; 9 T; 0 U; 0 Other;
Sequence 18
                                                                                                                                The present sequence was used in the development of a method
                                                                                                                                                                                                                                     WPI; 1997-558135/51.
                                                                                                                                                                                                                                                                                                                                                                     27-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                 11-NOV-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                              US5686242-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT96107;
                                                                                                                                                            Disclosure; Col 27-28; 22pp; English.
                                                                                                                                                                                                                                                                 Lima WF,
                                                                                                                                                                                                                                                                                              (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4299 CATCTTTTTCCTTCCCCT 4316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                              SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                Bruice TW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CATCTTATTCCTTTCCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
 BP; 1
                                                                                                                                                                                                                                                                                              PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer
                                                                                                                                                                                                                                                                                                                         91US-00755485.
92WO-US007489.
                                                                                                                                                                                                                                                                                                                                                                     94US-00330000
 Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.2%;
 0
 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
 G; 14 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1
Pred. No. 1.8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     research;
                                                                                      separating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                   ဓ္က
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR;
                                                            The
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BXXXE
                                                        RESULT 2921
AAX88679
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AAX63294/c
                                                                                                                                                        Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                      The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
                                                                                                                                                                                                                                                                                                                                                                                                             Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in
10-SEP-1999
                                                                                                                                                                                                      Sequence 18
                                                                                                                                                                                                                                                                                                                                                                           Claim 40; Page 93; 155pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-202224/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zwick MG,
Young SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modulation; gene expression; transgenic plant; cleavage; canola caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maize;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Delta-9 desaturase hairpin ribozyme substrate SEQ ID NO:1169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX63294 standard;
                                             AAX88679 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          granule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9710328-A2
                                                                                                           18
                                                                                                                                65
                                                                                                                                                        16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   corn; Zea mays; delta-9 desaturase; GBSS; target; substrate;
                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOWELANCO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bound
                                                                                                                                                                                                                                                                                                                                                                                                   canola.
                                                                                                        GCTGCTGCGGCGGCGGCG 1
                                                                                                                                ecreceeeeececece
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGATGTTTTTTTTTTT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Edington BE, Folkerts O,
                                                                                                                                                                                                      B₽;
                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          starch synthase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95US-0001135P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96WO-US011689
                                                                                                                                                                                                      2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA;
                                                                                                                                                                 0.2%;
                                                                                                                                                                                                      10 C;
                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Merlo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen JA,
Merlo DJ;
                                              ВP
                                                                                                                                                                                                      σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
                                                                                                                                82
                                                                                                                                                       0;
                                                                                                                                                    Score 14.8; DB 1; Length 18; Pred. No. 1.8e+03; 0; Mismatches 2; Indels
                                                                                                                                                                                                      G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hammerhead ribozyme;
                                                                                                                                                                                                      0
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U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Merlo PAO,
                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            canola plant;
                                                                                                                                                        ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Skokut TA;
                                                                                                                                                        Gaps
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                                                                                                                                                       0,
RESULT 2922
AAZ35875
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modulation; expression; diagnosis;

Human 03-FEB-2000 AAZ35875 standard; DNA; 18

ВP

0

AAZ35875

(first entry)

Human; sentrin; antisense oligonucleotide; phosphorothioate;

inhibition;

sentrin phosphorothioate antisense oligonucleotide SEQ ID NO:17.

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                                                                                                                                                                                          CC disorder or related disorder. AXX88542 to AAX88705 represents human CC chromosome 18q VAC clones and primers corresponding to them, used in the CC exemplification of the present invention. YAC clones comprising a portion CC of the region of human chromosome 18q between markers D1858 and D185979 CC are used to identify at least one human gene associated with a mood CC disorder or related disorder. The mood disorder or related disorder, is CC chosen from the Diagnostic and Statistical Manual of Mental Disorders, is CC version 4 (DSM-IV) taxonomy. This includes mood disorders (296, XX, 300.4, CC version 4 (DSM-IV) taxonomy. This includes mood disorders (295, XX, 300.4, CC version 4 (DSM-IV) taxonomy. This includes mood disorders (295, XX, 300.4, CC version 4 (DSM-IV) axisity disorders (300.XX, 309.81, 308.3), CC version the disorders (309.XX) and personality disorders (codes 301.XX). CC version the disorders (309.XX) and personality disorders (codes 301.XX). CC version the used to detect pathological mutations or genetic code contact of the susceptibility of an individual to a mood disorder or can be used to treat mood disorder. The muchoic acids and proteins of the human gene can be contact to the susceptibility of an individual to a mood disorder or can be contact to the treat mood disorders and related disorders.
                                                                             Matches
                                                                                                Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes detecting nucleotide triplet repeats region of human chromosome 18g disposed between polymorphic markers D18588 and D18599 to identify a human gene associated with a mood disorder or related disorder. AAX88542 to AAX88705 represents human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human chromosome 18q; mood disorder; polymorphic marker; detection; identification; trinucleotide repeat expansion; schizophrenia; anxiety disorder; adjustment disorder; personality disorder; nucleotide triplet repeat; ss.
                                                                                                                                                          Sequence 18 BP; 2 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 56; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUL-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human chromosome 18q YAC clone primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting nucleotide triplet repeats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-418934/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Van Broeckhoven C, Raeymaekers P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-DEC-1997; ·
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9932643-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                     5795 CCTGCCTGCCTGTC 5812
                                                                           16;
μ
                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VLAAMS
CCTGCCTACCAGCCTGTC
                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97GB-00026804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INTERUNIVERSITAIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-EP008543
                                                                                                  88.9%;
                                                                                                                     0.2%;
18
                                                                           0;
                                                                                                                   Score 14.8;
                                                                                                  Pred. No. 1.8e+03
                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INST BIOTECHNOG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Del-Favero J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human chromosome
                                                                                                                   DB 1; Length 18;
                                                                             2
                                                                             0
                                                                             Gaps
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
Homo sapie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes an antisense compound (I) 8-30 nucleotides long targeted to a nucleic acid molecule encoding human sentrin. The antisense compound comprises a phosphorothicate antisense oligonucleotide which inhibits expression of human sentrin. (I) is useful for inhibiting expression of sentrin in human cells or tissues in vitro, for treating humans or other animals suspected of having or being prone to a disease associated with sentrin expression. (I) can also be used for
                                                                                                                    vascular endothelial growth factor; VEGF; interleukin-10; IL-10; cancer; prostaglandin E2; PGE2; immune response; tumour; asthma; Crohn's disease; monocyte chemotactic protein-1; MCP-1; ulcerative colitis; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to a disease associated with sentrin expression. (I) can also be used for research or diagnostic purposes. The present sequence represents a human sentrin phosphorothicate antisense oligonuclectide from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
              16-DEC-1999.
                                       WO9963975-A2
                                                                    Unidentified
                                                                                               atherosclerosis.
                                                                                                            glomerulonephritis; acute
                                                                                                                                                                 Immunosuppressant inhibitor; transforming
                                                                                                                                                                                          Immunosuppressant inhibitor oligonucleotide TGF-betal-98-17.
                                                                                                                                                                                                                       30-MAR-2000
                                                                                                                                                                                                                                                                            AAZ65529 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense compound which modulates human sentrin expression, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-022284/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                       328

    Col 38; 29pp; English.

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                                                                                                                                                                                                                                                                                                                                               μ
                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                                                                                                                                 h 0.2%;
Similarity 88.9%;
16; Conservative
                                                                                                                                                                                                                                                                                                                                            CTGTCCAATGACTTTGAG 18
                                                                                                                                                                                                                                                                                                                                                                     CTGGCCAATTACTTTGAG 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diseases associated with sentrin expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 4 A;
                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-00213768
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 C; 4 G; 6 T;
                                                                                                            respiratory
                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1;
Pred. No. 1.8e+03;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 U;
                                                                                                            distress
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 18;
                                                                                                                                                 growth factor beta; TGF beta; interleukin-10; IL-10; cancer
                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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AAZ59187/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                  AAZ59187 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 1 A; 6 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 10; Fig 1; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Composition containing immune stimulant and inhibitor of agent that adversely affects the immune response, for treating cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-097470/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUN-1998;
25-JUL-1998;
                                                                   30-MAR-1999;
                                                                                                                                                                                                                 Fusion protein; TEV protease; P(
                                                                                                                                                                                                                                                                     Reverse primer for construct MWPsp-MWPmp5 DNA.
                                                                                                                                                                                                                                                                                                           15-SEP-2003
20-APR-2000
                                                                                                                                                                                                                                                                                                                                                                AAZ59187;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schlingensiepen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JUN-1999;
                                 31-MAR-1998;
                                                                                                                                             JP11341991-A
                                                                                                                                                                               Brevibacillus brevis
(ITOH-) ITOHAM FOODS INC
                                                                                                       14-DEC-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CTGGCAGCTGGCCGGGCG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                 n; Bacillus; cell wall protein; promoter; cleavage site; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98EP-00110709.
98EP-00113974.
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Pred. No. 1.8e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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RESULT 2925
AAZ48498
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Best Local S
Matches 16
The invention provides antisense compounds targeted to human tumour necrosis factor receptor type 1 (TWFR1) RNA. These antisense compounds can be used in a method of inhibiting the expression of TWFR1 human cel or tissues. The antisense compounds specifically hybridize with one or more nucleic acids encoding TWFR1 modulating the function of nucleic ac molecules encoding TWFR1, ultimately modulating the amount of TWFR1 produced. The antisense compounds and method are useful as research
                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sato
                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a DNA construct encoding a fusion protein comprising a Bacillus species cell wall protein fused to a cleavage peptide and a heterologous protein. The fusion construct is placed downstream of a Bacillus species promoter sequence. This sequence represents a PCR primer for the MPRBP-MPRBS part of the construct MPRBP-MPRBS-Met-ProInsulin, which comprises the Bacillus brevis middle wall protein mp5 linked to the human proinsulin protein. (Updated on 15-SEP-
                                                                                       Example
                                                                                                                      Antisense
                                                                                                                                                                                                            26-JUN-1998;
                                                                                                                                                                                                                                 26-JUN-1998;
                                                                                                                                                                                                                                                        28-DEC-1999.
                                                                                                                                                                                                                                                                                                                                               Tumour necrosis
                                                                                                                                                                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                                                                                                                                                                         31-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                AAZ48498;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A DNA
                                                                                                                                                                                                                                                                                                                                    inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UDAK/)
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                                                                                                          inhibition, treatment
                                                                                                                                                                  Cowsert
                                                                                      Col 24; 34pp;
                                                                                                                                                                                                                                                                                                                                                                   mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression.
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                                                                                                                                                                                                            9803-00106038
                                                                                                                                                                                                                                  98US-00106038
                                                                                                                                                                                                                                                                                                                                  factor receptor type 1; TNFR1; antisense;
umour formation; TNFR1; anticancer; ss.
                                                                                                                                                                                                                                                                                                                                                                   inhibiting
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                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                          of tumor necrosis and prevention of
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No. 1.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preparation
                                                                                                        factor type 1 expression for disease, particularly tumors.
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AAZ71698/c
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                               AAZ6554 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical efficacious responses to and side effects from differential sefificacious responses to and side effects from the Dharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID Nos 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches 16
                                                                                                                                                                                                                                     Claim
 Sequence
                                                                                                                                                                                                                                                               map
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23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; identification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        reagents and diagnostics, and in the treatment and prophylaxis infection, inflammation or tumour formation. Sequences AAZ48482 represent antisense oligos used for inhibition of the human TWF
                                                                                                                                                                                                                                                                             Novel biallelic markers used
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                                                                                                                                                                                                                                     Page 1521;
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                                                                                                                                                                                                                                                                                                                                 Blumenfeld
                                                                                                                                                                                                                                                              human genome
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98US-0109732P.
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Pred. No. 1.8e+03;
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  Other;
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Query Match Best Local

Similarity

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14.8; DB 1; No. 1.8e+03;

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RESULT 2927
AAZ76847/c
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Best Local
                                                                                                                                               agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a semicone in the figure of pharmaceutical agents.
                                                                                                                                                                                                              invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-2001
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                                                                                                       Sequence 18
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23-NOV-1998;
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                                                                                                                                   present invention
                                                                                                                                                                                                                                                                                                                           AAZ65654 to AAZ69578 represent human biallelic markers from the present
                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 2619; 2745pp; English
                                                                                                                                                                                                                                                                                                                                                                                            Novel biallelic markers
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98US-0109732P.
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RESULT 2928

Homo sapiens

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RESULT 2929
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Best Local S
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                                                                                                                                                                                                                                                                                                 chemical or drug in a sample comprises contacting the sample with an oligonucleotide having specific activity for a target biomolecule and detecting the presence or absence of binding where the presence of binding indicates the presence of the chemical or drug in the sample. The oligonucleotide is identified by: (a) assaying a prepared set of randomized oligonucleotides for activity against a target biomolecule; (b) separating active from inactive oligonucleotides; and (d) characterizing the recovered oligonucleotides; and (d) characterizing the recovered can be used for diagnostic or research nurroses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-SEP-1991;
04-SEP-1992;
27-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ88678 standard;
                                        Human; anti-inflammatory; cytostatic; antisense inhibition; inflammation; tr
                                                                                                                                       AAA30403 standard; DNA; 18
                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      an
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Assay for a chemical or drug in a sample comprises de
an oligonucleotide selected from a set of randomized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-170669/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6022691-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chimeric primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ88678;
                               cancer; 88.
                                                                       Human NF-kappa-B
                                                                                             21-AUG-2000
                                                                                                                  AAA30403;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide selected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WF,
                                                                                                                                                                                                              4460 GGACTTTTTTTTTTTT
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                                                                                                                                                                                                                                     l Similarity
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                                                                                                                                                                                                                                                                              BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                 Col 27-28; 20pp; English.
                                                                                                                                                                                                                                      Conservative
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                                                                                            (first entry)
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92WO-US007489.
94US-00330000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-00965908
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                                                                        p65 subunit antisense oligodeoxynucleotide
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Pred. No. 1
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                                                                                                                                                                                                                                      Mismatches
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                                          transcription factor; apoptosis;
                                                     antimicrobial; infection
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                                                                                                                                                                                                                                                 1.8e+03;
                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                0 Other;
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                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      etecting binding of oligonucleotides.
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                                                                          ISIS# 23770.
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RESULT 2930
AAA38376
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                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense compounds which inhibit the expression of the human NF-kappa-B p65 subunit (p65) useful for treating diseases associated with p65 expression and as prophylaxis to prevent of delay infection, inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                          Ets-2; human; transcription factor; chromosome 21q22.3; cancer; metastasis; skeletal abnormality; Down's syndrome; expression in
                                                                                                                  Human Ets-2 phosphorothioate antisense oligonucleotide, SEQ ID NO:35
                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAY-2000
            US6054316-A
                                      Homo sapiens
                                                                phosphorothicate;
                                                                                                                                                 21-AUG-2000
                                                                                                                                                                                                   AAA38376
                                                                                                                                                                                                                                                                                                                                                                                                          automated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or tumor formation.
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                                                                                                                                                                                                                                                                                               2124
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                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                   standard;
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                                                                                                                                                                                                                                                                                                                                                                               18 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                        DNA synthesiser
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                                                                                                                                               (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                antisense;
                                                                                                                                                                                                   DNA; 18
                                                                                                                                              entry)
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Pred. No. 1.
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on, inflammation
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                                                                            r; invasion; inhibition;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC a DNA-binding domain shared by all members of this family. Through this cC motif, Ets family members bind to the promoter regions of various genes CC at a GCA consensus sequence, thereby acting as either repressors or activators of the gene. All but one Ets family protein bind to DNA as a CC monomer. Ets-2 has been implicated in the regulation of cellular CC proliferation and differentiation. The Ets-2 gene is located at CC chromosome 21q22.3, which is within a region known to undergo CC translocations associated with malignancies. Ets-2 has been found to be C upregulated in several cancers, including lymphoblastic leukaemia. It may CC also play a role in the cancer phenotype, as it activates the urokinase c plasminogen activator (uPA) promoter and the promoters of metalloproteinases in response to epidermal growth factor (EGF) C metalloproteinases in response to opidermal growth factor (EGF) C with tumour invasion and metastasis in breast cancers. As the Ets-2 gene is located on chromosome 21, which is triplicated in Down's syndrome, it is also thought to be responsible for the skeletal abnormalities present in this condition. The antisense oligonucleotides of the invention are useful for the treatment or prophylaxis of conditions associated with Ets
                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequences AAA38349-A38388 represent antisense oligonucleotides targetted to the human Ets-2 gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human Ets-2 RNA, and were analysed for their effect on Ets-2 mRNA levels by quantitative real-time PCR. The Ets-domain transcription factors are a family of proteins which are involved in controlling key cellular events such as proliferation, differentiation and development. The Ets domain is
                                                                                                                                                       S. noursei
                                                                                                  Polyketide synthase; PKS; macrolide; nystatin; PKS gene cluster; antifungal; antibiotic; enoylreductase; ER; PCR primer; ss.
                                                                                                                                                                                         29-NOV-2001
                                                                                                                                                                                                                           AAD17190;
                                                                                                                                                                                                                                                      190/c
AAD17190 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense compound, 8-30 nucleobases Ets-2 is useful for treating cancer a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Col 39; 31pp; English
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                                                                                                                                                                                                                                                                                                                                                                                7231 ATCCCTCTCAAGTCCAGC 7248
                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                     PKS-encoding NysC DNA amplifying PCR antisense primer
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                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 18;
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16-AUG-2001

WO200159126-A2 Streptomyces noursei

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ARESULT 2932
AAH4604
AC AAH4604
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CA SYNTHE
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Matches 16; Conserv
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Best Local
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10-APR-2000;
14-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to the cloning and sequencing of the gene cluster encoding a modular type I polyketide synthase (PKS) enzyme involved in the biosynthesis of the macrolide antibiotic nystatin. The nystatin PKS is useful as antifungal antibiotics. The present sequence a PCR primer which is used for the amplification of the DNA fragment representing the coding sequence for the C-terminal part of the encylreductase (ER) domain in module 5 of Streptomyces noursei PKS-
                                                                                                                                                                                                                                                             Synthetic oligonucleotide; dinucleotide repeat; cell cycle arrest; cell proliferation; caspase; tumour necrosis factor; TNF; cancer; carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 4 A;
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(SNTF )
(ALPH-)
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       13-DEC-1999;
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(ZOTC/)
(SEKU/)
(FJAE/)
                                                                                                                                                                                                                                                                                                                                                          Synthetic oligonucleotide 23.
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                                                12-DEC-2000; 2000WO-CA001467.
                                                                                                                                               WO200144465-A2
                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                       lymphoma; ss.
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S, Ellingsen
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ZOTCHEV S B.
SEKUROVA O N.
FJAERVIK E.
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STROM A R.
VALLA S.
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llingsen TE,
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2000GB-00009387.
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                                                                                                                                                                                                                                                                                                                                                                                                           (first entry
    99US-0170325P
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Pred. No. 1.
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                                                                                                                                                                                                                                                             cytostatic; apoptosis;
cytokine; interleukin;
sarcoma; leukemia;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Composition comprising synthetic oligonucleotides which comprise multiple repeats of dinucleotides such as GT, TG useful for treating cancer by inducing cell cycle arrest, inhibiting proliferation, activating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 19; Page 32; 77pp; English.
                                                                                                                                                                                                                                                                          Synthetic.
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                                                                                                                                   24-JAN-2002.
                                                                                                                                                                                                                            misc_RNA
                                                                                                                                                                                                                                                                                                      DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                   DNA-RNA-DNA oligonucleotide AGT02013 used to test RNase H cleavage.
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                                                                                                                                                                                                                                                                                                        detection;
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TECHNOLOGIES INC

14-JUL-2000; 30-MAR-2001;

2000US-00616761 2001US-00823647

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ABL40838
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotides required for RNase H cleavage. Each oligonucleotide of the set had a different number of ribonucleotides, 2 in the present case. The oligonucleotides were mixed with target DNA oligonucleotide AGT02009 (see ABA91531) and incubated with RNase H (5 U/ml) at 37 degrees C for 30 minutes. The results showed that 4 ribonucleotides were the minimum number for RNA cleavage. The invention provides probes for nucleic acid hybridisation. The probes form a hairpin structure comprising a double-stranded stem and a single-stranded loop, and are capable of both intramolecular and intermolecular hybridisation. The double-stranded stem may comprise a methylphosphonate DNA:RNA hybrid that is resistant to RNase H cleavage. When the probe hybridises with a target DNA, the RNA strand in the DNA:RNA duplex becomes sensitive to RNase H treatment and can be removed. Arrays and methods for nucleic acid hybridisation using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           exbB; exbD; tonB; antibiotic; toluene; pHBA; aromatic compound; parabene; para-hydroxybenzoic acid; RT-PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                 30-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                      28-SEP-2001; 2001WO-US031180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pseudomonas putida
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                                                                                                                                                                                                                                                                                                                                                                                                                                                11-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                exbD genes amplifying RT-PCR primer.
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Disclosure; Page 80; 81pp; English

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Matches 16; Conserv
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       conditions from tissue samples. The method comprises obtaining a sample from test tissue or region to be diagnosed, performing a methylation assay of the sample, where the assay determines methylation state of genomic CpG sequences, and making a diagnostic or prognostic prediction of the cancer based at least in part upon the methylation state of the genomic CpG sequences. The method is useful for diagnosing cancer or cancer-related conditions such as gastrointestinal or esophageal adenocarcinoma, gastrointestinal or esophageal dysplasia, gastrointestinal or esophageal metaplasia, Barrett's intestinal tissue, pre-cancerous conditions in normal esophageal squamous mucosa, or their combinations, from tissue samples. Preferably, the cancer diagnosed is esophageal adenocarcinoma, and making a diagnostic or prognostic
                                                                                                                                                                                                                                                                                         Diagnosing cancer or cancer-related conditions, e.g., gastrointestinal and esophageal adenocarcinoma, comprises performing methylation assay a tissue sample obtained from a test tissue or region to be diagnosed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel gene cluster comprising the exbB, exbD and tonB genes from P. putida. These genes are useful for producing bacterial cells more sensitive to antibiotics, toluene, pHBA (parahydroxybenzoic acid), aromatic compounds, parabenes, and aromatic amino acids. Methods are also provided to identify pHBA tolerant genes, and pHBA tolerant strains, useful for producing pHBA. The present sequence represents a primer for RT-PCR amplification of P. putida exbB and exbD
                                                                                                                                                                                                                      The invention relates to a method of diagnosing cancer or cancer-related
                                                                                                                                                                                                                                                          Claim 3; Page 32; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                  WPI;
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Pred. No. 1.8e+03;
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RESULT 2936
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             The present invention describes a method of arraying genome clones. The CC method comprises: (a) clones of the genomic libraries contained in CC multiwell plates numbered for discrimination are mixed in each of the CC multiwell plates; (b) a primer designed based on the chromosome marker gequence is added to the mixture to carry out an amplification reaction; (CC (c) a signal corresponding to the marker is detected from the resultant CC amplified product to specify the discrimination Nos. of the multiwell CC plates containing the clones having said marker sequence; (d) the order CC of the markers is changed so that the seme discrimination Nos. succeed to the markers is changed so that the seme discrimination Nos. succeed to CC the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified CC discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified by using the above primer; (g) signals are detected from the amplified producte solt; and (1) the clones are reconstituted as the positions on the chromosome and arrayed. The mixed respects and arrayed the primers of the producte of the positions of the chromosome and arrayed. The chromosome are constituted as the positions on the chromosome and arrayed.
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Matches 16; Conserv
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1p36-35 DNA,
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and ABL45323
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Best Local :
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                                                                                                                                                                                                                       The invention relates to an antisense compound 8 to 30 nucleotides in length targeted to nucleic acid molecule encoding tumour necrosis factor receptor 1 (TNFR1), where the antisense compound inhibits expression of TNFR1. The antisense compound is useful for inhibiting the expression of TNFR1 in cells or tissues. The antisense compound is also useful for treating an animal (preferably human) having a disease or condition associated with TNFR1, e.g. a liver disease (such as hepatitis, or liver injury) or a hyperproliferative disorder such as cancer, by inhibiting the expression of TNFR1. The antisense compound is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. This property of the expression of the expression of TNFR1 antisense compound is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel antisense compound targeted to nucleic acid molecule encoding necrosis factor receptor 1 (TNFR1), useful for treating humans having disease associated with TNFR1 e.g. hepatitis, liver injury, liver continues.
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                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 having
/er cancer.
                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                ABZ11028
                                                               RESULT 2939
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ABX80015
                                                                                                                                                                             Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                        repeat within a coding sequence (expressed sequence tag, EST), which comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenaemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying a candidate polymorphic repeat within a coding sequence, understanding or treating genetic disease, comprises detecting tande repeats in a target coding sequence and scoring the repeats for polymorphic probability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPO Rep-X; human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome;
ABZ11028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST polymorphic DNA repeat polynucleotide #340.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABX80015 standard; cDNA; 18 BP
                              ABZ11028 standard; DNA; 18
                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention discloses a method for identifying a candidate polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 spinal atrophy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fredreich's ataxis; myotonic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABX80015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                7415
                                                                                                               \mathbf{L}
                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                               18
                                                                                                                                              GCAGCAGCAGCAGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Col 1165; 588pp; English.
                                                                                                                                                                                                                                               B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wren JD,
                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-00475947.
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                                                                                                                                                                                                                                               5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cis; myotonic dystrophy; hyperandrogenaemia;
bulbar atrophy; spinocerebellar ataxia.
                                                                                                                                                                                             0.2%;
                                                                                                                                                                                                                                             8 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Minna JD,
                                BP
                                                                                                                                             7432
                                                                                                               18
                                                                                                                                                                             0;
                                                                                                                                                                            Score 14.8; D
Pred. No. 1.8e
0; Mismatches
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                                                                                                                                                                                              1.8e+03
                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                            Length 18;
                                                                                                                                                                             Indels
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ABZ11030/ ID ABZ1 XX

ABZ11030 standard; DNA; 18

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RESULT 2940

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                                                                                                                      Best Loc
Matches
                                                                                       Query Match
                                                                                                                                            lymphocytic leukaemia and acute myelogenous leukaemia; as probes for determining the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) of haematopoietic cell proliferation disorder related sequences and their complements; and as primers for the amplification of haematopoietic cell proliferation disorder related DNA sequences. The nucleotide sequences from the present invention can also be used for detecting a predisposition to, differentiation between subclasses, diagnosis, prognosis, treatment and/or monitoring of haematopoietic cell proliferative disorders. The present method enables highly specific classification of haematopoietic cell proliferative disorders allowing for improved and informed treatment of patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Berlin K,
Olek A,
Lewin A,
                                                                                                                                                                                                                                                                                                                   differentiating between haematopoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent, which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid. ABZ09861 to ABZ11118 represent specifically claimed nucleotide sequences from the present invention. Oligonucleotides from the present contact and proliferative disorder haematopoietic cells; for differentiating between acute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy; lymphocytic leukaemia; acute myelogenous cytosine methylation state; probe; primer; ss.
                                                                                                                      Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2001; 2001US-0278333P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Haematopoietic cell proliferation disorder related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2002; 2002WO-EP003401.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EPIG-)
                                                                       Local
                              3210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003-018942/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              K, Braun A, Distler J, Guetig D, Piepenbrock C, Adorjan P, Grabs A, Lipscher E, Maier S, Model F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             haematopoietic cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EPIGENOMICS
                                                                       Similarity
                            TGAGAAAGTGGGTGGGAG 3227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lipscher E, I
                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         77; 117pp;
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                                                                      0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
18
                                                         0;
                                                                                       Score 14.8; DB 1;
                                                                                                                    G; 4 T; 0 U; 0 Other;
                                                                       Pred.
                                                            Mismatches
                                                                       No.
                                                                         .8e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for detecting and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Howe A,
Lesche R
                                                                                      Length 18;
                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide #1168.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mueller
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ر ق:
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RESULT 2941
ABZ81757
ID ABZ8175
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                                                                                                                                                                                CC differentiating between hasmatopoietic cell proliferative disorders cassociated with at least 1 gene and/or their regulatory regions in a cc subject. The method comprises contacting a target nucleic acid in a cc biological sample obtained from the subject with at least 1 reagent, cc which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid. ABZ09861 to ABZ1118 cc represent specifically claimed nucleotide sequences from the present cc invention. Oligonucleotides from the present invention can be used: for cd differentiating between healthy hasmatopoietic cells and proliferative cc lasorder hasmatopoietic cells; for differentiating between acute myelogenous leukaemia; as probes for cd determining the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) of hasmatopoietic cell proliferation disorder related sequences and their complements; and as primers for the cc related sequences and their complements; and as primers for the cc sequences. The nucleotide sequences from the gresent invention can also be used for detecting a predisposition to, differentiation between complements, treatment and/or monitoring of hasmatopoietic cell proliferative disorders. The present method enables a highly specific classification of haematopoietic cell proliferative disorders. The present method enables a disorders allowing for improved and informed treatment of patients.
                                                                                                               Matches
                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
Synthetic.
 ABZ81757 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; Page 77; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-018942/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-2001; 2001US-0278333P
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                                                                                                                                                               Sequence 18 BP; 4 A; 11 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPIG-) EPIGENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention describes a method for detecting and
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éI,
                                                              18
                                                                                                               16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        K, Braun A, Distler J,
, Piepenbrock C, Adorjan
A, Lipscher E, Maier S,
                                                                                                                            Similarity
                                                                                        TGAGAAAGTGGGTGGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lipscher E,
Ziebarth H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ą
                                                                                                                      0.2%;
                                                                                      3227
                                                                                                              0,
                                                                                                                           Score 14.8;
Pred. No. 1.
                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Guetig D, Howe A, M
P, Grabs G, Lesche
Model F, Mueller V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disorder related
                                                                                                                           1.8e+03
                                                                                                                                      DB 1;
                                                                                                                                     Length 18;
                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Otto T,
                                                                                                              0,
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                                                                                                               Gaps
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RESULT 2942
ADA27361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene. Preferred oligonucleotides are single-stranded, have at least one mismatch with respect to the HD gene region to be altered, and have chemical modifications, or are chimeric RNA/DNA oligonucleotides. They can be used for the treatment or prevention of HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-AUG-2001; 2001US-0310757P.
08-AUG-2001; 2001US-0310770P.
08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
                                                                                   ds; HLA-related research; HLA class II-associated disease; transplantation matching; recombination hot spot identification; linkage disequilibrium study; human; microsatellite.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is an example of a poly-glutamine triplet repeat region found in exon 1 of the Huntington's disease (HD) gene. The invention is based on the discovery that oligonuclectides can be designed to target sequence alterations to the triplet repeat region of the HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New single stranded oligonucleotides comprising a DNA domain least one mismatch with respect to the genetic sequence of th Huntington's disease gene to be altered, useful for treating preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-AUG-2002; 2002WO-US025352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Huntington's disease; nootropic; anticonvulsant; huntingtin; human; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Huntington's
US2003108940-A1
                                                                                                                                                                             Human microsatellite repeat M2_3_8.
                                                                                                                                                                                                                                                                            ADA27361;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 57; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-256478/25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                          Homo sapiens
                                                                                                                                                                                                                               20-NOV-2003
                                                                                                                                                                                                                                                                                                                          ADA27361 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7407 CAACATCAGCAGCAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                         CAACAACAGCAGCAACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease
                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 6 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                          DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
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                                                                                                                                                                                                                                                                                                                          BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                         18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14.8; DB 1;
No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18;
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ADB54474
ID ADB5
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an oligonucleotide primer capable of specifically hybridising to a DNA having the sequence of the flanking regions of a microsatellite selected from M2-4-9, M2-2-9, M2-2-12, M2-3-11, M2-2-20, M2-2-21, M2-2-22, M2-2-23, M2-2-24, M2-4-25, M2-4-26, M2-2-29, M2-2-32, M2-4-32, M2-4-33, M2-4-37, M2-3-32, M2-2-36, M2-5-11, M2-2-46, and M2-2-48. The primer is useful for determining the number of repeat units of the microsatellite cited above. The primer is useful in HLA-related research, such as genetic mapping of HLA class II-associated disease, transplantation matching, population genetics, and identification of recombination hot spots as well as linkage disequilibrium studies. The present sequence represents the human microsatellite repeat M2_3_8.
                      Adorjan I
Rujan T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligonucleotide primer capable of specifically hybridizing to a DNA having the sequence of the flanking regions of a microsatellite (e.g. M249), useful for HLA-related research, e.g. transplantation matching.
WPI; 2003-731620/69
                                                                                                                              04-SEP-2003
                                                                                                                                                                                                              colon cell;
cytostatic;
                                                                                                                                                                                                                                                 Hybridisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-DEC-2002;
                                                                               27-FEB-2002;
                                                                                                       27-FEB-2003; 2003WO-EP002035
                                                                                                                                                      WO2003072821-A2
                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                        04-DEC-2003
                                                                                                                                                                                                                                                                                               ADB54474;
                                                                                                                                                                                                                                                                                                                     ADB54474 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-NOV-2000; 2000US-00713616.
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                                                        (EPIG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
16; Conserv
                    P, Burger M,
, Schmitt A;
                                                        EPIGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                       GCTGCGGGGGGGCGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tamiya G,
                                                                                                                                                                                                                        proliferative disorder; non methylated CpG dinucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 5; 20pp; English.
                                                                                                                                                                                                              cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 0 A;
                                                                               2002EP-00004551
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                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                               oligonucleotide 12
                                                                                                                                                                                                                                                                                                                     DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
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                                  Maier
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                                                                                                                                                                                                                                                                                                                                                                                                    82
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Pred. No. 1.
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                                                                                                                                                                                                             carcinoma; cytosine
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                                 Nimmrich I,
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                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8e+03;
                                                                                                                                                                                                                                              analyse
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 Other;
                                 Becker
                                                                                                                                                                                                             methylation state;
                                                                                                                                                                                                                                              genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 18;
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Best Local S
Matches 16
                                                                                                                                 Burger M,
Nimmrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel method for detecting and differentiating between colon cell proliferative disorders associated with at least one gene or its regulatory regions. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least one reagent or a series of reagents, where the reagent or series of reagents, distinguishes between methylated and non methylated CpG dinucleotides within the target nucleic acid. The molecules of the invention demonstrate cytostatic activity whilst the method may useful for detecting and differentiating between colon cell proliferative disorders, including cancers such as colon adenoma and colon carcinoma. The PNA (peptide nucleic acid)-oligomers are useful as probes for determining cytosine methylation state or single nucleotide polymorphisms. The current sequence is that of the hybridisation oligonucleotide of the invention which was used to analyse the genomic
                                                                                                                                                                                                                           10-DEC-2002; 2002WO-EP014026
                                                                                                                                                                                                                                                                                WO2003052135-A2
                                                                                                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                                                                                                  cytosine methylation state.
                                                                                                                                                                                                                                                                                                                                              PCR; primer; ss; lung cell proliferative adenocarcinoma; squamous cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                 18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 36; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting and differentiating between colon cell proliferative disorders associated with a gene or its regulatory regions comprises contacting a target nucleic acid in a biological sample obtained from the subject with
                                                                                                                                                                        (EPIG-) EPIGENOMICS
                                                                                                                                                                                                 14-DEC-2001; 2001DE-01061625
                                                                                                                                                                                                                                                     26-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                                       Primer oligo used
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC69952;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC69952 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6672 TTGGGGGACGTTATTTTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTGGGGGATGTTATTGTT 18
                                                                                                                                             Field
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                             Ę,
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                                                                                                                                                                                                                                                                                                                                                                                     for analysing CpG islands in genomic DNA (SeqID 441).
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                                                                                                                                             Genc
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                                                                                                                                            В,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1;
Pred. No. 1.8e+03;
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                                                                                                                                            Liloglou
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                                                                                                                                            'n
                                                                                                                                                                                                                                                                                                                                                cytostatic;
                                                                                                                                                                                                                                                                                                                                                            disorder; CpG dinucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 Other;
                                                                                                                                             Lipscher
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                                                                                                                                                                                                                                                                                                                                              probe;
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                                                                                                                                                                                                                                                                                                                                                PNA-oligomer;
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Detecting and differentiating cytosine methylation state of genomic DNA, useful for diagnosing, treating prognosticating and/or monitoring lung cell proliferative disorders e.g. adenocarcinoma and squamous cell

WPI; 2003-533029/50

Claim 15;

SEQ

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441;

58pp;

English

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RESULT 2945
ADE43413/c
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a novel method for detecting and differentiating between lung cell proliferative disorders associated with at least one gene and/or their regulatory regions. Specifically, it refers to a method comprising contacting a target nucleic acid in a biological sample with at least one reagent, wherein the reagent is able to distinguish between methylated and non-methylated Cpd dinucleotides present in the target DNA. As such, it is possible to further adifferentiate and diagnose medical conditions including adenocarcinoma and squamous cell carcinoma, and their respective adjacent lung tissue. The present invention describes cytostatic oligomers and PNA-oligomers that are useful as probes for determining the cytosine methylation state or single nucleotide polymorphisms (SNPs) of the target sequence. This oligomucleotide sequence is a primer oligomer used for the analysis of CpG positions within genomic DNA, used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                     Neurodegenerative disease; uPA; SNCG; IDE; KNSL1; LIPA; TNFRSF6; Alzheimer's disease; neuroprotective; nootropic; gene therapy; Chromosome 10; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human SNCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18
                                                                          Determining a predisposition for or the occurrence of neurodegenerative disease, e.g. Alzheimer's disease by detecting in a target nucleic acid the presence or absence of an allelic variant of one or more polymorphic
                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE43413;
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                                                                                                                                                                                                                                                                   08-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE43413 standard; DNA; 18
                                                                                                                            WPI; 2003-559131/52.
                                                                                                                                                     Saunders
                                                                                                                                                                 Becker KD,
                                                                                                                                                                                                                                                      09-NOV-2001;
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                                                                                                                                                                                                                                                                                             25-OCT-2001;
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                                                                                                                                                                                         NEUROGENETICS INC. GEN HOSPITAL CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequencing primer, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
ilarity 88.9%;
Conservative
                                                                                                                                                                                                                             ; 2001US-0339525P.
; 2001US-0336929P.
; 2001US-0338010P.
; 2001US-0338363P.
; 2001US-0337052P.
; 2002US-0368919P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                                   Velicelebi
                                                                                                                                                                                                                                                                                                                      2002WO-US034679.
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                                                                                                                                                     Mullin
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                                                                                                                                                    Elliott
Sampson A
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Pred. No. 1.8e
O; Mismatches
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Blacker I
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The present invention relates to a method (M1) for determining a predisposition for or the occurrence of neurodegenerative disease

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6716 CAGGATGGTAGCTGGAAT 6733

Conservative

0; Mismatches

Indels

0

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0

2

Page

265;

848pp; English

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                                                       RESULT 2946
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Best Local 9
Best Local Similarity
Matches 16; Conserv
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              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        subject. The method comprises detecting in a target nucleic acid obtained from the subject the presence or absence of an allelic variant of one or more polymorphic regions of one or more genes selected from upA (Urokinase plasminogen activator), SNCG (gamma-symuclein), IDE (insulindegrading enzyme), KNSL1 (Kinesin-like protein 1), LIPA (lysosomal acid lypase), and TNFRSF6 (Tumour Necrosis Factor Receptor-SF6), where the presence of at least one of the allelic variant of one or more polymorphic regions is indicative of a predisposition for or the occurrence of neurodegenerative disease. The genes are all located on chromosome 10. M1 is useful for determining a predisposition for or the occurrence of, and for treating neurodegenerative disease, particularly Alsheimer's disease. The present sequence is a PCR primer, which was used in the method of the invariance.
                                                                                                                                                            Assay reagent contg. oligo-nucleotide probe attached via probe having hybridisation region complementary to specif sequence, e.g. for diagnosis of genetic disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe fixed via 400nt poly-dT linker to a nylon Beta-thalassemia allele.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-OCT-2002
14-MAY-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 5 A; 7 C; 4 G; 2 T; 0 U;
                                                                               probe fixed to a filter allows simultaneous non radioactive detection 50 or more specific nucleotide sequences in a single test sample. (Updated on 31-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                Saiki RK,
                                                                                                                                                                                                                                                                                                                                                                         30-NOV-1989.
                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAN92724;
                                                       Sequence 19
                                                                                                                                                                                                                      WPI; 1989-370739/50.
                                                                                                                                                                                                                                                                                                      04-MAY-1989;
                                                                                                                                                                                                                                                                                                                   20-MAY-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe; fixed; oligo-nucleotide; hybridisation;
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                                                                                                                                                                                                                                                                            (CETU ) CETUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the method of the invention.
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                                                                                                                                     Page 29; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTTCTCTCTGGGGAGGGA
                                                                                                                                                                                                                                                 Erlich HA;
                                                       BP;
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(first en
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89US-00347495.
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                                                       5 A;
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C;
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               Pred.
                           Score 14.8;
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              No.
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               .9e+03
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                             DB 1;
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                           Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  identify target
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RESULT 2947
AAQ25298/c
ID AAQ2529
  RESULT 2948
AAQ49843/c
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AC AAQ4984
XT 25-MAR-
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KW Human I
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The oligomer is a target sequence for complexing with a synthetic oligomer to inhibit transcription. The oligomer binds to the transcribed region of the DNA so as to form a triple helix. The target region may like within an exon or an intron and the oligomer forms a triple helix by expliciting the CT motif (i.e. the oligomer is pyrimidine rich). The oligomer is useful in antisense therapy. It can also be used (opt. in labelled form) diagnostically to detect target DNA or RNA by hybridisation. See also AAQ25290-300. (Updated on 25-MAR-2003 to correct the property of the contract of th
Human Immunodeficiency Virus; antigen; ELISA; recombinant antibody;
                                                                                                                                                                                                        AAQ49843 standard; cDNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Table 1; Page 26; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inhibiting transcription of duplex DNA in diagnosis - by contacting the transcribed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Transcription;
triple helix;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Target sequence for ODN5.
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                                            PCR primer S2 to amplify signal region
                                                                                              25-MAR-2003
27-APR-1994
                                                                                                                                                                AAQ49843;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1992-234645/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-DEC-1990;
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23-DEC-1992
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(first entry)
                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; inhibition; riplex; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     triplex;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90US-00625680.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        duplex; anti-sense therapy; RNA; polymerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽P
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Pred. No. 1.9e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     1.9e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               anti-sense therapy and region of DNA with an oligomer
                                                                                                                                                                                                                                                                                                                                                                                                                2
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AAQ53087/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EBV-transformed cell lines and mouse-human heterohybridomas producing human MAbs specific for the gp120 V3 loop of HIV-1 MN isolate were obtained. MAb 447-52D was found to recognise the tetrappetide motif GPGR, i.e. the Principal Neutralising Determinant common to the V3 loop of different HIV isolates. A recombinant Ab was derived from 447-52D comprising the H and L chain V regions, each fused to a signal sequence and to appropriate intronic sequences. The different fragments required to construct the fusion sequences were amplified by PCR. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV-neutralising monoclonal antibody; immunoglobulin; AIDS; acquired immune deficiency syndrome; chimeric antibody; surface glycoprotein gp120; V3 loop; ss.
                                   13-FEB-1992;
                                                           10-SEP-1992;
                                                                                                             JP05285000-A.
                                                                                                                                     Synthetic
                                                                                                                                                                       Gene detection; radio-isotopes; target gene; electrode; detection;
optical_fibre; hybridise; hybridisation; electrochemical; photochemical;
                                                                                                                                                                                                                                      03-JUN-1994
                                                                                                                                                                                                                                                                                      AAQ53087 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 9; Fig 3; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New recombinant human antibody - with HIV neutralising activity against at least two isolates, useful for preventing or treating infection in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Emini EA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-APR-1992;
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                                                                                    02-NOV-1993.
                                                                                                                                                                                                                                                              AAQ53087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MERI ) MERCK & CO INC.
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                                                                                                                                                                                                                                                                                                                                                                            6022 TCCACACCTGTCCACTCC 6039
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                                                                                                                                                                                                              detection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 4 A; 3 C; 9 G; 3 T;
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                            probe;
                                    92JP-00025621
                                                             92JP-00242397
                                                                                                                                                                                                            sequence
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                              1.9e+03
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RESULT 2950
AAA85905
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequences (AAQ53077-Q53136) are used in the invention to detect specific genes without the use of radio-isotopes. Detection is carried out by hybridisation of denatured (ss) sample DNA with a (ss) nucleic acid probe, complementary to the target sequence. Hybridisation occurs on the surface of an electrode or optical fibre and detection is visualised by the addition of an entity that recognises (ds) hybridised DNA and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection method of gene without using radio-isotope - by hybridisation of nucleic acid probe which is single strand having complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1993-382240/48
                                The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                     04-DEC-1998;
                                                                                                                                                                                                                                                                                                              06-DEC-1999;
                                                                                                                                                                                                                                                                                                                                      08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                             WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                            Ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Cdc 25 hs ribozyme
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Sequence
                                                                                                                                    Disclosure;
                                                                                                                                                                       New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle
                         restenosis
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                                                                                                                                                           Cyclin
                                                                                                                                                                                                                                    Welch PJ,
                                                                                                                                                                                                                                                                                                                                                                                                            hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid probe which is single strand having gene and single strand denatured sample
                                                                                                                                   Page 99; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 3 A;
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 B₽;
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3
A;
                                                                                                                                                                                                                                                             INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     binding site
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8 C; 1
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Pred. No. 1
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 T, 0
                                                                                                                                                                                                                                      Robbins
                                                                                                                                                                        for inhibiting restenosis, cleaves dependent kinase other than CDK1,
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 Other;
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Query Match

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Score 14.8;

ВB

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Length

19;

Ribozyme; hairpin;

hammerhead; gene therapy; vasotropic; restenosis;

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                                                                                                                                                                                                 RESULT 2951
AAA82490/c
EXEXEXEX
                                                                                          RESULT 2952
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                                                                                                                                                             Matches
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Best Local
                                                                                                                                                                                                                          designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1.
Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA82490 standard;
                                                                                                                                                                                                                                                                                                                          New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle PCNA and Cyclin B1.
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                                    04-DEC-2000
                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                      Disclosure; Page 47; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-412314/35
                                                                                                                                                                                                                                                                                                                                                                                  Tritz
                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cdk1 ribozyme binding site
                Cdc 25
                                                       AAA85904;
                                                                         AAA85904 standard;
                                                                                                                                                                                                                                                                                                                                                                                                     (-DWMI)
                                                                                                                                                                                                                                                                                     present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1215 TACCTACCTTCCCCTAGA
                                                                                                                                          6736 CTTCCTTCTTAAAATCTG 6753
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                                                                                                                                                                                                  19 BP;
                                                                                                                         CTTCCTTTTTAGAATCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hairpin;
                 ribozyme
                                                                                                                                                                                                                                                                                                                                                                                  Welch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                      invention relates to a hairpin or hammerhead ribozyme
                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0110954P
                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-US028772
                                                                                                                                                                                                                                                                                                                                                                                                     INC.
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                                                                                                                                                                                                                                                                                                                                                                                  PJ,
                                                                                                                                                                                                  Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hammerhead;
                binding site #12.
                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                    0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                   Barber
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                                                                                                                                                                               Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene therapy; vasotropic; restenosis;
                                                                                                                                                                                                  w
                                                                                                                                                             Mismatches
                                                                                                                                                                                                   1
                                                                                                                                                                                                                                                                                                                                                                                   Robbins
                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                   for inhibiting restenosis, cleaves dependent kinase other than CDK1,
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                                                                                                                                                                       .9e+03
                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                   Ϋ́
                                                                                                                                                                                DB 1;
                                                                                                                                                                               Length 19;
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                                                                                                                                                              Indels
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                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                               kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88
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RESULT 2953
AAZ71461
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin BI. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                 Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotrype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                     28-OCT-1999
                                                                                                                                                                                                                                        Human biallelic marker upstream amplification primer SEQ ID NO:5817.
                                                                                                                                                                                                                                                                                                              AAZ71461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New hairpin and hammerhead RNA encoding a cyclin or ce PCNA and Cyclin B1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-JUN-2000.
 Cohen D,
                                             21-APR-1998;
23-NOV-1998;
                                                                                21-APR-1999;
                                                                                                                            WO9954500-A2
                                                                                                                                                 Homo sapiens
                                                                                                                                                                        diagnosis; ss.
                                                                                                                                                                                                                                                                 10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 99; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-DEC-1998;
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                       (GEST )
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16; Conserv
                      GENSET
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                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                             TACCTACCTTCCCCTAGA 1232
                                                                                                                                                                                                                                                                                                                                                                        TACCTCCTTTCCCCCTAGA
 Blumenfeld
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                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                (first entry)
                                             98US-0109732P
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                                                                                99WO-IB000822.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ΡJ,
                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
Σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Barber JR,
                                                                                                                                                                                                                                                                                                               19
 Chumakov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ad ribozyme
cell-cycle
                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Robbins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for inhibiting restenosis, cleaves dependent kinase other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                              1.9e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
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AAC65640/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                             Detecting and analyzing genotypes of polymorphic loci amplified in a mixture of amplified restriction fragments, particularly amplification fragment length polymorphism fragments using primer extension techniques
                     This invention describes a novel method for determining polymorphic loci amplified in a restriction fragment mix
                                                                                                                                                                                      Kuiper MTR,
                                                                                                                                                                                                                                                09-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                             Human; AFLP; pamplification
                                                                                                                                                                                                                                                                                                                                                                                                                                          Human AFLP primer E51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC65640 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 ar 3367, are not actually given a sequence in the Sequence Listing from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       map of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel biallelic markers used to construct a high density disequilibrium
                                                                   Example 1; Page 17;
                                                                                                                                                                                                                                                                            10-APR-2000; 2000WO-NL000235.
                                                                                                                                                                                                                                                                                                         19-OCT-2000
                                                                                                                                                                                                                                                                                                                                    WO200061801-A2
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                                                                                                                                                                                                                   (KEYG-)
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                                                                                                                                                          2000-679499/66
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                                                                                                                                                                                                                                                                                                                                                                                                            AFLP;
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                                                                                                                                                                                                                   KEYGENE NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCTTCCTTTTCCTCTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 1471;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCCTTTCTTTTCCTTTTC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                      Witsenboer
                                                                                                                                                                                                                                                                                                                                                                                             polymorphic loci; SNP; single nucleotide polymorphism;
a fragment length polymorphism; genetic marker; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7 C; 1
                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5719
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1;
Pred. No. 1.9e+03;
a restriction fragment mixture (M), usin S) complementary to part of target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
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polymorphic loci an oligonucleotic

oligonucleotide sequence (OS)

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RESULT 2955
AAS42915
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                           23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
02-MAR-2000;
03-MAR-2000;
03-MAR-2000;
13-MAR-2000;
03-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fragment length polymorphism (AFLP) fragments such that useful non-polymorphic bands, which ordinarily do not provide any useful information when conventional AFLP-finger printing is used, are made. SNPs are also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  metabolīc disorder; Parkinson's disease; Tourette's syndrome; thrombosi type 2 diabetes; cardiovascular disorder; myocardial infarction; cancer cardiomyopathy; atherosclerosis; human immunodeficiency virus; HIV; viral infection; immunostimulant; neuroleptic; nootropic; tranquiliser; antidepressant; anorectic; PCR primer; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS42915;
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                                                                                                                                                                                                                                                              23-JUN-2000;
11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200162797-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        attention deficit disorder; anxiety; depression; bipolar disord neurological disorder; Huntington's disease; dementia; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human G Protein-Coupled Receptor (GPCR) PCR primer #49.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                informative as genetic markers
                                               WPI; 2001-570628/64.
                                                                                                                                              (PHAA)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G-protein coupled receptor; GPCR; mental disorder;
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                                                                                                                                           PHARMACIA & UPJOHN CO.
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2000US-0184304P.
2000US-0184305P.
2000US-0184305P.
2000US-0186457P.
2000US-0186810P.
2000US-018880P.
2000US-018880P.
2000US-0194344P.
2000US-0213861P.
2000US-0213879P.
2000US-0213837P.
2000US-0218337P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001WO-US005676.
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                                                                                              Parodi
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            schizophrenia;
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thrombosis;
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New isolated nucleic acid

encoding a new G-protein coupled receptor

The present invention describes an improved method of DNA synthesis provides double stranded DNA where the predetermined orientation of sequence is preserved. This can be used in the construction of complementary DNA and directional DNA libraries

which the

Double stranded DNA synthesis with specific orientation comprises synthesizing a first strand of DNA complementary to a selected DN template and synthesizing second strand complementary to first on

DNA one

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WPI; 2001-006435/01.

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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA molecules encoding human G-protein coupled receptor (GPCR) polypeptides. The protein and DNA sequences of the invention can be used to identify compounds which bind to GPCR polypeptides and in screening for compounds that modulate GPCR activity. By screening a human subject for the presence of mutations in GPCR DNA, a GPCR-related disorder or a genetic predisposition can be diagnosed. The sequences can also be used for treatment and prevention of mental disorders such as schizophrenia, attention deficit disorder, anxiety, depression, dementia and bipolar disorder, neurological disorders such as Huntington's disease and Tourette's syndrome, metabolic disorders such as obesity, anorexia and type 2 diabetes, cardiovascular disorders such as thrombosis, myocardial infarction, cardiomyopathy and atherosclerosis, viral infections caused by HIV and cancers
                                                                                                                                                                                                                            19-SEP-1988;
02-MAY-1991;
23-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences AAS42806-AAS42926 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptide for detecting receptor modulators that can treat mental disorders, such as schizophrenia, anxiety, depression, or obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                      DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC83562 standard; DNA; 19 BP
                                                                                                                                                                                                                  02-SEP-1993;
                                                                                                                                                                                                                                                                              22-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                           DNA synthesis;
                                                                                                                                                                                         (STRA-)
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                                                                                                                                                                                                                                                                                                                                                                                                                    synthesis
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                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                               88US-00246567.
91US-00700066.
92US-00981931.
93US-00116049.
                                                                                                                                                                                                                                                                                                                                                                                                                    method linker/primer sequence SEQ ID NO:
                                                                                                                                                                                                                                                                              97US-00899029
                                                                                                                                                                                                                                                                                                                                                                                             directional complementary DNA library; linker;
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No. 1
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ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, antisickling, cytostatic, antiseborrheic, antidiabetic, antisickling, cytostatic, antiseborrheic, antidiabetic, antisickling, cytostatic, antiseborrheic, antidiabetic, antisickling, cytostatic, antiseborrheic, antidiabetic, antisickling, cophthalmological, cytostatic and cytocid in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local
Matches
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                                                                                                                                                                                                                                                                                                                                                             Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipporiatic; dermatological; antiseborrheic; antidiabetic; virucide; antipical; antiseborrheic; antidiabetic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                     The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sickle cell retinopathy; ss.
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                                                                                                                                                                                                                                                                                                                             Page 325; 408pp; English.
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Pred. No. 1.
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Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                     dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                exemplification of the present invention Sequence 19 BP; 3 A; 8 C; 1 G; 7 T; 0 U; 0 Other;
diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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                                                                                                                                                                                                                                                                                                                                              The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-300427/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic l scar. AAH57577 to AAH62099 represent sequences used in the exemplification of the present invention
The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
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16; Conserv
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Pred. No. 1.
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RESULT 2960
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PAX2;
                                                                                                                                                                                                      Modulating resistance to apoptosis, rescuing cells from apoptosis, enhancing resistance of normal tissues to apoptotic cell death induced chemo- or radiation therapy in patients by using PAX-2 function
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24-JUL-2000; 2000US-0220161P.
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tion therapy; cancer; prostate; ovary; bladder; kidney;
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CC single strand conformation polymorphism (RT-PCR-SSCP) of PAX2 exon 8.

CC PAX2 is a transcription factor involved in the development of the nervous CC and excretory systems and mutations of PAX2 have been associated with CC optic nerve colobomas and renal hyperplasia. These mutations are CC modulating resistance to apoptosis. The method of the invention involves CC modulating resistance of normal tissues to apoptotic cell death induced by CC chemotherapy or radiation therapy. This is achieved by administering to a CC patient, a nucleic acid (I) encoding a molecule which selectively CC inhibits and/or prevents the function of the PAX2 gene. The method can be CC used to modulate resistance to apoptosis of cancer cells (prostate, cystic kidney cells in a patient in which PAX2 is expressed at higher CC cystic kidney cells in a patient, for rescuing cells from apoptosis or level than in a healthy patient, for rescuing cells from apoptosis in a

Disclosure; Page

23; 45pp; English

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ABL88900
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Matches
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                                                                                              The present invention describes a collection of binding groups for a family of nucleic acids comprising members of relative high and relative low significance, where the binding groups are selected to be capable to identify, alone or in combination, essentially all members of the family of nucleic acids of relatively high significance. The collection of binding groups is useful for typing of nucleic acid in a clinical sample, by contacting the nucleic acid with the collection and determining whether one or more binding groups bound to the nucleic acid of the sample. This method is useful for determining whether the sample comprises at least a part of a member of relatively high significance of a family of nucleic acids. The collection of binding groups is useful for diagnosing the severity of a disease caused by a pathogen containing a member of a family of nucleic acids. ALBSF79 to ALBSF31 represent
                                                                                                                                                                                                                                                                                                                     Collection of binding groups for determining or typing samples, especially clinical samples, has groups capable to identify ess all members of the family of nucleic acids of relatively high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                Disclosure; Page 36; 166pp; English.
                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-156696/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUL-2000; 2000EP-00202611.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Binding
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                                                  Sequence
                                                                                         oligonucleotide
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16; Conserv
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Pred. No. 1.9e
0; Mismatches
Score 14.8; DB 1,
Pred. No. 1.9e+03,
0; Mismatches :
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RESULT 2963
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ABA97625/c
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Best Local S
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      Screening; G protein-coupled receptor; cholesterol metabolism; inflammatory disease; transplantation rejection; immune insuff infection; PCR; primer; TNF alpha.
                                                                                                                                                                                                                                    This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
                                                                                                                                                                                                                                                                                                               Measurement
the obtained
                                           TNF alpha
                                                                               AAL51775;
                                                                                                AAL51775 standard; DNA; 19
                                                                                                                                                                                                                    Sequence
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(KANK-)
(KEIZ-)
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24-AUG-1999; 99JP-0023666.
30-AUG-1999; 99JP-00242693.
01-FEB-2000; 2000JP-00028896.
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KEIZAI SANGYOSHO SANGYO
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                                            PCR
                                                                                                                                                                                                                                                                                             Page 17;
                                                                                                                                                                                                                                                                                                               of nucleic acids,
d data.
                                                                                                                                                                                                                    BP;
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                                                             (first entry
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                                                                                                                                                                                                                    P.
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No. 1.9e+03;
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RESULT 2964
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Screening method for agonists or antagonists to alter binding properties of novel G protein-coupled receptor protein in controlling cholesterol metabolism, used to diagnose and treat inflammatory diseases or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-APR-2001; 2001JP-00114203.
14-JUN-2001; 2001JP-00180565.
16-JUL-2001; 2001JP-00214922.
27-DEC-2001; 2001JP-00397767.
22-FEB-2002; 2002JP-00045728.
                                           11-FEB-2003; 2003WO-US004034
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                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                           RNA interference; cytostatic; vasotropic; nephrotropic; modulation; inhibition; cancer; breast cancer; ovarian cancer; lung cancer; prostate cancer; glioblastoma; proliferative disease; restenosis; polycystic kidney disease; human; ribozyme; ss.
                                                                                                                        28-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      short interfering nucleic acid; siNA; protein kinase C alpha;
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16; Conserv
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.9e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                  The present invention describes a short interfering nucleic acid (siNA) CC that downregulates expression of a protein kinase C-alpha (PKC-alpha) CC gene by RNA interference. Also described: (1) a siNA that modulates CC expression and/or activity of genes for other isoforms of PKC or genes CC involved in the PKC pathway; (2) kits for in vitro or in vivo delivery of SiNA; (3) conjugates and/or complexes of siNA; and (4) vectors that CC express siNA. The siNA sequences have cytostatic, vasotropic and CC express siNA. The siNA sequences have cytostatic, vasotropic and CC of expression of the PKC-alpha gene by RNA interference. The siNA can be useful in treating a variety of cancers including e.g. breast cancer, CC cancers of the head and neck, ovarian cancer, lung cancer, prostate CC cancers of the head and neck, ovarian cancer, lung cancer, prostate CC cancers of the head and for treating other proliferative diseases CC and conditions, such as restenosis and polycystic kidney disease. The SiNA may also be useful for diagnosis, drug screening, target condition, and for gene mapping (e.g. of single-nucleotide polymorphisms). CC The present sequence represents a human PKC-alpha siNA target, which is CC used in the exemplification of the present invention.
                                       Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0366782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
18-SEP-2002; 2002US-0411707P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis of cancer and restenosis, protein kinase C-alpha gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 118; 143pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New short interfering nucleic acid,
                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-679891/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SIRN-)
       3937 TCCCTTGATGGTCAAGTC
                                        l Similarity
16; Conserv
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                                                                                                          B₽;
                                       0.2%;
llarity 88.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Beigelman
                                                                                                          6
                                                                                                          A; 4 C; 5
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        3954
                                        0;
                                                                                                          G; 0 T; 4 U;
                                                                         Score
                                                          Pred.
                                          Mismatches
                                            3 14.8; _______;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            useful e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            downregulates
                                                                                                          0 Other;
                                                                          ۲.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ġ
                                          <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for treatment
                                                                       Length 19;
                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression
                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of the
                                          Gaps
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RESULT 2965
ADA25417
short interfering nucleic acid; siNA; protein kinase C alpha; PKC-alpha; RNA interference; cytostatic; vasotropic; nephrotropic; modulation; inhibition; cancer; breast cancer; ovarian cancer; lung cancer;
                                                                                                                                                                     Human PKC-alpha short interfering nucleic acid SEQ ID NO:148
                                                                                  polycystic kidney
                                                                                                                                                                                                 20-NOV-2003
                                                                                                                                                                                                                              ADA25417;
                                                                                                                                                                                                                                                          ADA25417 standard; RNA; 19
                                                                                               prostate cancer; glioblastoma; proliferative disease;
                                                                                                           nhibition; cancer; breast cancer; ovarian cancer;
                                                                                                                                                                                                 (first entry)
                                                                                  disease; human; ribozyme;
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WO2003070983-A1

domo sapiens.

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                                                                                                                                                                                                                                          RESULT 2966
                                                                                                                                                                                                                                                                                                                                                                                                                                CC The present invention describes a short interfering nucleic acid (sina) CC that downregulates expression of a protein kinase C-alpha (PKC-alpha) CC gene by RNA interference. Also described: (1) a sinA that modulates CC expression and/or activity of genes for other isoforms of PKC or genes CC involved in the PKC pathway; (2) kits for in vitro or in vivo delivery of CC sinA; (3) conjugates and/or complexes of sinA; and (4) vectors that CC express sinA. The sinA sequences have cytostatic, vasotropic and CC of expression of the PKC-alpha gene by RNA interference. The sinA can be useful in treating a variety of cancers including e.g. breast cancer, CC cancers of the head and neck, ovarian cancer, lung cancer, prostate cancer, and glioblastoma and for treating other proliferative disease. The sinA may also be useful for diagnosis, drug screening, target conditions, such as restenosis and polycystic kidney disease. The condition, and for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a human PKC-alpha sinA, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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09-SEP-2002;
18-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
                                        stearoyl-CoA desaturase; RNA interference; anorectic; antidiabetic; antiarteriosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening;
                            genetic engineering;
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein kinase C-alpha gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen
                                                                                      short interfering nucleic acid; siNA; downregulation; inhibition;
                                                                                                                     Stearoyl-CoA
                                                                                                                                                   29-JAN-2004
                                                                                                                                                                                                             ADE27307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-FEB-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                     3937 TCCCTTGATGGTCAAGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2003-679891/64.
                                                                                                                                                                                                                                                                                                                                                 11;
                                                                                                                                                                                                                                                                                       H
                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3; Page 118; 143pp; English.
                                                                                                                                                                                                            standard; RNA; 19
                                                                                                                                                                                                                                                                                       UCCCUUGAUGGAUAAGUC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         interfering nucleic acid, useful e.g. for treatment and of cancer and restenosis, downregulates expression of the cancer and restenosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; 2002US-0363124P.
; 2002US-0386782P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0409293P.
; 2002US-0411707P.
                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                    desaturase sinh oligonucleotide SEQ ID NO:251.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2003WO-US004034.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2003US-0440129P.
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                                                                                                                                                  (first
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                                                                                                                                                                                                                                                                                                                                                                                                          A,
                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                          5 C; 4 G; 0 T; 6 U; 0 Other;
                            pharmacogenomic;
                                                                                                                                                                                                                                                                                       18
                                                                                                                                                                                                                                                                                                                  3954
                                                                                                                                                                                                                                                                                                                                              5
                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1;
Pred. No. 1.9e+03;
5; Mismatches 2
                            gene mapping;
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                                                                                                                                                                                                                                                                                                                                                                          Length 19;
                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                      SCD;
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ADE27597/c
ID ADE2759
XX
ADE2759
XX
AC ADE2759
XX
AC ADE2759
XX
BC Stearoy
XX
Short i
XW short i
XW stearoy
XW stearoy
XW stearoy
XW stearos
XW antiart
XW antiart
XW antheriox
XX
Synthet
XX
XX
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                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a short interfering nucleic acid (siNA) CC that downregulates expression of the SCD (stearcyl-CoA desaturase) gene CC by RNA interference. Also described: (1) modulating expression of SCD CC genes in cells, tissue explants or organisms by introduction of siNA; (2) conjugates in vitro or in vivo delivery of siNA; (3) conjugates and/or CC complexes of siNA; and (4) vectors that express siNA. SCD inhibiting CC siNAs have anorectic, antidiabetic, antiarceriosclerotic, cytostatic and CC virucide activities. The siNAs can be used to modulate expression of SCD CC genes, in cells, tissue explants or organisms, e.g. for treating obesity; CC diabetes (types I and II); atherosclerosis; cancer and viral infections. CC identification and validation; genetic engineering; pharmacogenomics; cudying gene function and gene mapping (e.g. of single-nucleotide cc polymorphisms). The present sequence represents an SCD siNA, which is constructed the complification of the present invention.
                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
20-SEP-2002;
                                                        antiarteriosclerotic; cytostatic; virucide; obesity; diabetes;
atherosclerosis; cancer; viral infection; drug screening;
                                                                                    short interfering nucleic acid; siNA; downregulation; stearoyl-CoA desaturase; RNA interference; anorectic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New short interfering nucleic acid, useful e.g. diagnosis of obesity or diabetes, downregulates stearoyl-CoA desaturase gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003070885-A2
                                         genetic engineering;
                                                                                                                                  Stearoyl-CoA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2002;
11-MAR-2002;
                                                                                                                                                              29-JAN-2004
                                                                                                                                                                                             ADE27597
                                                                                                                                                                                                                        ADE27597 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JAN-2003;
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                                                                                                                                                                                                                                                                                                                            4471 TTTTTTTTTTTTGTCTT 4488
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; 2002US-0358580P.
; 2002US-0363124P.
; 2002US-0366782P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0409293P.
; 2002US-0412304P.
; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                 desaturase sinh oligonucleotide SEQ ID NO:541.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003WO-US004317
                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ID NO 251; 139pp;
                                                                                                                                                                                                                                                                                                                                                                                                                 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM
                                                                                                                                                                                                                                                                                                                                                                    5.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                 0 C;
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                                                                                                                                                                                                                                                                                                                                                                      Pred.
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                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                   0 T;
                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                       No.
                                                                                                                                                                                                                                                                                                                                                                                                                 15 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                      .9e+03;
                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for treatment expression of
                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                     inhibition;
                                                                                       antidiabetic;
                                                                                                                                                                                                                                                                                                                                                          .
                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                     SCD;
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Synthetic

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                                                                                                                                                                                                                                                              RESULT 2968
AAN81957/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a short interfering nucleic acid (siNA) CC that downregulates expression of the SCD (stearcyl-CoA desaturase) gene by RNA interference. Also described: (1) modulating expression of SCD CC genes in cells, tissue explants or organisms by introduction of siNA; (2) CC kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or CC complexes of siNA; and (4) vectors that express siNA. SCD inhibiting CC siNAs have anorectic, antidiabetic, antiarteriosclerotic, cytostatic and virucide activities. The siNAs can be used to modulate expression of SCD CC genes, in cells, tissue explants or organisms, e.g. for treating obesity; CC diabetes (types I and II); atherosclerosis; cancer and viral infections. CC identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide CC polymorphisms). The present sequence represents an SCD siNA, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0366782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
20-SEP-2002; 2002US-0412304P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid, useful e.g. for treatment and diagnosis of obesity or diabetes, downregulates expression of the stearoyl-CoA desaturase gene.
15-DEC-1988
                                                                                                                                                                                                                                            .957/c
AAN81957 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 541; 139pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-FEB-2003; 2003WO-US004317.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-AUG-2003
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                              WO8809820-A.
                                                                                                       Lymphocyte function associated antigen-3; adhesion inhibition; probe;
                                                                                                                                      Probe pool LF1 to detect Human Lymphocyte Function Associated Antigen-3
                                                                                                                                                                    25-MAR-2003
22-OCT-1990
                                                                                                                                                                                                                 AAN81957;
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                                                                                                                                                                                                                                                                                                                                                     4471
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                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 15
                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                   (revised)
(first en
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                                                                                           immune suppression;
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                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                             0 G; 0 T; 1 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                    0;
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Pred. No. 1.9e+03;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                 Length 19;
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RESULT 2969
AAQ03650
ID AAQ0365
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BIOJ )
(DAND )
(DAND )
(DAND )
                                                                                                                                                                                                                                   PI-linked lymphocyte function associated antigen 3 polypeptide; phage lambda P24; T-cells; autoimmune disease; graft versus host disease; allograft rejection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA sequences encoding Lymphocyte Function Associated Antigen-3 - inhibits adhesion between T-lymphocytes and target cells.
                                                                                                                                                                                                                                                                                                             25-MAR-2003
07-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                  AAQ03650 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 1 A; 3 C; 1 G; 10 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page ?; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1988-368634/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wallner BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-JUN-1987;
                             New DNA sequences and recombinant DNA - expressing PI-linked lymphocyte function associated antigen-3 polypeptide.
                                                                                                                            26-AUG-1988;
                                                                                                                                                  24-AUG-1989;
                                                                                                                                                                        08-MAR-1990.
                                                                                                                                                                                           WO9002181-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-JUN-1988;
                                                             WPI; 1990-099405/13
                                                                                  Wallner BP,
                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                        Probe NN-D for use in Church-Gilbert sequencing of cDNA encoding
                                                                                                        (BIOJ ) BIOGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                           6344 ААСАТАЛАСССБААСААССТ 6363
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity 65.0
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DANA FARBER CANCER
DANA FARBER CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BIOGEN NV
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                                                                                  Hession
                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
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Pred. No. 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length
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Disclosure; Fig

5; 34pp; English

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RESULT 2971
AAQ03651/c
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AAQ03648/c
                                                                                     Query Match
Best Local S
Matches 16
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Probe NN-D for was used along with 3 other 20-nucleotide long probes, and NotI digestion, for sequencing by Church-Gilbert approach, of the ends of an inserted sequence in pNN01. This plasmid is a subclone of P24 contg. cDNA encoding PI-linked LFA-3. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                Probe NN-B for was used along with 3 other 20-nucleotide long probes, and NotI digestion, for sequencing by Church-Gilbert approach, of the ends of an inserted sequence in pNN01. This plasmid is a subclone of P24 contg. CDNA encoding PI-linked LFA-3. (Updated on 25-MAR-2003 to correct PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PI-linked lymphocyte function associated antigen 3 polypeptide; phage lambda P24; T-cells; autoimmune disease; graft versus host allograft rejection; ss.
                                                                                                                                  Sequence
                                                                                                                                                                                                                      Disclosure; Fig 5; 34pp; English.
                                                                                                                                                                                                                                           New DNA sequences and recombinant DNA - expressing PI-linked lymphocyte function associated antigen-3 polypeptide.
                                                                                                                                                                                                                                                                                                   Wallner BP, Hession
                                                                                                                                                                                                                                                                                                                                                                                                                 WO9002181-A.
                                                                                                                                                                                                                                                                                                                                                26-AUG-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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07-AUG-1990
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                                                                                                                                                                                                                                                                                                                          (BIOJ ) BIOGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linked LFA-3.
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                                                                 7307
                                                                                                                                                                                                                                                                               1990-099405/13.
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                                                                                       l Similarity
16; Conserv
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                                                               CTTTGAGATTTGTGTTTG 7324
                                                                                                                                  20 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTTTGAGATTGGTGGTTG 18
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                                                                                       Conservative
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                                                                                              0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
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                                                                                                                               8 C; 1 G; 3 T; 0 U;
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                                                                                                Score 14.8;
Pred. No. 26
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Pred. No. 2
                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8; DB 1;
2e+03;
                                                                                                2e+03;
                                                                                                                                  0 Other;
                                                                                                          DB 1;
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                                                                                                          Length
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                                                                                       Indels
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                                                                                     Gaps
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ID AAQ14994 standard;
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                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                               Query Match
                           04-MAY-1990;
                                                                                   Synthetic.
                                                                                                    human immunodeficiency virus; phosphorothioate linkage; retrovirus; ribosomal frame shift; gag; pol; fusion protein; ss.
                                                                                                                                Oligonucleotide #9 for modulating HIV-1 gag/pol frameshifting
                                                                                                                                                    24-FEB-1992
                                                                                                                                                                      AAQ14994;
                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                    Probe LF-1 is a 32-fold degenerate 20-mer derived from the LFA-3 purified from human erythrocytes. The probe was used libraries of PBL CDWA for clones encoding PI-linked LFA-3.
-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                               New
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-AUG-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PI-linked lymphocyte function associated antigen 3 polypeptide; phage lambda P24; T-cells; autoimmune disease; graft versus host disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
07-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ03651 standard; DNA; 20
                                             14-NOV-1991
                                                                WO9117246-A.
                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 5; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                     function
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wallner
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       allograft rejection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Probe LF-1 for cDNA encoding PI-linked LFA-3 (3'-5').
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ03651;
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                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                              DNA
                                                                                                                                                                                                                                                          6344 AACATAAAGCCGAAGAAGGT 6363
                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                    sequences and recombinant DNA - n associated antigen-3 polypeptid
                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BIOGEN
                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                       AAYAGRAARACRAARAAGGT 1
                                                                                                                                                                                                                                                                                                                 BP; 1 A; 3 C; 1 G; 10 T; 0
                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hession
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(first en
                                                                                                                                                    (first
                            90US-00518929
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                                                                                                                                                                                        DNA;
                                                                                                                                                    entry)
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                                                                                                                                                                                        20
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                                                                                                                                                                                                                                                                                      Score 14.8;
Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                    polypeptide.
                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                   U; 5 Other;
                                                                                                                                                                                                                                                                                      2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                              expressing
                                                                                                                                                                                                                                                                                               DB 1;
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04-MAY-1990;

90US-00518929

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RESULT 2973
AAQ29648/c
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Query Match
Best Local S
Matches 16
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This oligonucleotide and its analogue, having phosphorothicate bonds, were designed to specifically bind to the gag-pol frameshift region and interfere with translation and/or frameshifting. There is potential for significant RNA secondary structure near the site of frameshifting in HIV 1. The inhibitory effect of the oligo and its analogue has not yet been
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOdulating gene expression for HIV treatment - comprises binding oligonucleotide(s) to RNA portions which have sec. structure.
                                                              This PCR primer was used with AAQ29645 to detect the presence Hepatitis C viral RNA in a sample. (Updated on 25-MAR-2003 to field.)
                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                    Non-A non-B hepatitis; NANBH; HCV; detection; diagnosis; primer; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
16-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ29648 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A;
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                                                                                                       Disclosure;
                                                                                                                            Detection of primers with
                                                                                                                                                                                 Okamoto
                                                                                                                                                                                                                                                                    28-OCT-1992
                                                                                                                                                                                                                                                                                         EP510952-A1
                                                                                                                                                                                                                                                                                                              Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                 PCR primer #55 for identifying Hepatitis C virus
                                          Sequence
                                                                                                                                                            WPI; 1992-359137/44.
                                                                                                                                                                                                                           26-APR-1991;
                                                                                                                                                                                                                                               23-APR-1992;
                                                                                                                                                                                                      (IMMO ) IMMUNO JAPAN
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Local Similarity hes 16; Conserv
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                                                                                                                                                                                                                                                                                                                                   polymerase chain
                                          20
                                                                                                        Page 36;
                                                                                                                                                                                Nakamura
                                         B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
 Conservative
                                                                                                                            non-A, non-B hepatitis virus - us nucleotide sequences corresp. to
                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
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                                                                                                      54pp;
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          0.2%;
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                                                                                                        English
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                                          G; 9
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Score 14.8; DB 1;
Pred. No. 2e+03;
0; Mismatches 2;
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Pred. No. 26
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                                          T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                          Other;
                                                                                                                            using new oligo-nucleotide to part. of the viral RNA.
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                     Length
 Indels
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                      20
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RESULT 2974
AAQ24460/c
ID AAQ2446
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                                                                                                                  AAQ31484/c
                                                                                                                            RESULT 2975
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                             The sequences given in AAQ24460 and AAQ24461 are PCR primers which are used to amplify the 5' region of non-A, non-B hepatitis virus (NAMBHV) strain HC-J5. These probes amplify the region corresponding to nucleotides 867-1354 of the entire nucleotide sequence and were used to produced clones C5164, C5303 and C5331. The nucleotide sequences derived from this amplification can be used to detect NAMBHV infection which could not be detected by conventional methods. The detection kits allow highly specific and sensitive detection at an early phase of infection
                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant cDNA of NANBH virus strain HC-J5 and corresp. peptides - useful for diagnosis and in vaccines and immunological pharmaceuticals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Non-A.
non-B hepatitis virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               non-A, non-B hepatitis virus; NANBHV; HC-J5; PCR; amplification polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ24460;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ24460 standard;
                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 7; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-160959/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAY-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP485209-A.
                    NANB hepatitis virus PCR primer #55.
                                           25-MAR-2003
02-APR-1993
                                                                               AAQ31484;
                                                                                                     AAQ31484 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IMMO ) IMMUNO JAPAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NANB hepatitis virus primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2943 AACAGGGCCAGCAAGACA 2960
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                                                                                                                                                                                                                       Similarity
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                                                                                                                                                               AGCAGGGCCAGCAAGAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGCAGGGCCAGCAAGAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nakamura
                                                                                                                                                                                                                                                         BP; 1 A; 6 C; 4 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                            Conservative
                                           (revised)
(first ent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91EP-00310297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                             entry)
                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INC.
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                                                                                                                                                                                                            Score 14.8; DB Pred. No. 2e+03; 0; Mismatches
                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                    Length
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Polymerase chain reaction; non-A non-B hepatitis; detection;

88

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THE SECOND SECON
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AAQ43865/c
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence is that of PCR primer #55 which was used to determine the 5' terminus sequence from nucleotides 867-1354 of non-A, non-B hepatitis (NANBH) virus strain HC-J5. These nucleotide sequences encode structural proteins of NANBH virus and these proteins can be analysed to locate and provide polypeptides useful as antigens for detection of NANBH virus via antibody-antigen complex detection. Mutants, variants or fragments of the sequence can be used for very sensitive detection. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Non-A, non-B; specific; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NANB hepatitis viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
Antigen related to non-A and non-B hepatitis virus - comprise translation region comprising 340 - 341 mols. of nucleotides,
                                                                              WPI; 1993-199637/25
                                                                                                                                                                                  12-JUN-1990;
08-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                               JP05091884-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-OCT-1993
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                                                                                                                                                                                                                                                           10-APR-1991;
                                                                                                                                                                                                                                                                                                             16-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-APR-1991;
                                                                                                                                (NAKA/) NAKAMURA
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    -A, non-B hepatitis virus related antigens, their polynucleotide(s)
antibodies - are useful for detecting NANBH virus in blood samples
ended for transfusion.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    non-B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 1 A; 6 C; 4 G; 9 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    virus;
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90JP-00304405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymerase chain reaction; detection; sensitive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene HC-OM PCR primer #55
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Pred. No. 26
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                            comprises
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Matches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 2977
         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                  AAQ74282 and AAQ74283 are the forward and reverse PCR primers for human amyloid precursor protein (ApP) exon 14, these were used to screen yeast artificial chromosome (YAC) libraries for APP. Isolated APP clones were then injected into blastocysts, from the same species as the embryonic cells which contained the YAC library. Transgenic animals which could be used as models of beta-amyloidosis (prevalent in individuals with Down's syndrome and Altzheimers disease), were then generated from the injected blastocysts. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification by PCR of numerocation by PCR of numerocation by PCR of numerocations with the patities virus gene HC-OM
                                              Sequence 20
                                                                                                                                                                            Example 1; Page 24; 60pp;
                                                                                                                                                                                                Introduction and expression of large genomic sequences in transgenic animals - which may be used as animal models of beta-amyloidosis in Alzheimer's disease and Down's syndrome.
                                                                                                                                                                                                                                                WPI; 1994-333207/41.
                                                                                                                                                                                                                                                                        Gearhart JD,
                                                                                                                                                                                                                                                                                                                                              01-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                           WO9423049-A2
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                       yeast artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amyloid precursor protein; APP; exon 14 PCR primer;
beta-amyloidosis animal models; Down's syndrome; Alzheimers disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ74282;
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                                                                                                                                                                                                                                                                                               (UYJO ) UNIV
                                                                                                                                                                                                                                                                                                                      02-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                     13-OCT-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amyloid precursor protein exon 14 forward PCR primer
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12-JUN-1995
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                                              BP; 4 A;
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                                                                                                                                                                                                                                                                                               JOHNS HOPKINS.
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(first en
                                                                                                                                                                                                                                                                        Lamb BT;
                                                                                                                                                                                                                                                                                                                      93US-00042390
                                                                                                                                                                                                                                                                                                                                              94WO-US003619
                                                                                                                                                                                                                                                                                                                                                                                                                                         chromosome; ss.
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                                              5 C; 5 G;
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                                                                                                                                                                            English.
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Score 14.8; I
Pred. No. 2e+0
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8; I
Pred. No. 2e+0
0; Mismatches
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                                              6 T;
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           2e+03;
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                                              0 Other;
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                       DB 1;
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                       Length
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                       20;
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standard; DNA;

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25-MAR-2003 19-OCT-1995 AAQ98015 AAQ98015

(revised) (first entry)

PNA oligomer

antiviral; antisense; Peptide nucleic acid;

triple

helix; ss.

PNA; HIV; human immunodeficiency virus; AIDS

HIV gag/pol frameshift.

/*tag=

Location/Qualifiers

moiety in the next residue"

RESULT 2978 AAQ98015/c

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29-JUL-1993;
                                                                                                                                                                                              28-JUL-1994;
                                                                                                                                                                                                         09-FEB-1995.
                                                                                                                                                                                                                   WO9504068-A1
                                                                                                                                                                                                                                                                       misc_feature
                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                 /note= "at least one (and preferably all) of the backbone subunits are composed of N-acetyl N-(2-aminoethyl)glycine peptide residues, the nucleosebe being stratched covalently to the acetyl group and the peptide linkage being formed by condensation of the glycine carboxy group of one residue with the amino group of the 2-aminoethyl
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New peptide nucleic acid (PNA) oligomers are provided which (a) consist of naturally occurring nucleobases covalently bound to a polyamide consist of hackbone and (b) hybridise to the translation initiation AUG region, 5 (cuntranslated region (5 UTR), 3 untranslated region (3 UTR), splice concerns or coding sequence of a human immunodeficiency virus gene concerns from env, gag, pol, rev and tat. The PNAs can be used to target RNA and single stranded DNA (ssDNA) to produce antisense-type gene concerns for the processes. Hence they can be used to treat AIDS and other viral infections. They are also useful in diagnostic applications and as createrch tools. PNA oligomers have high affinity for complementary single cresent translated DNA. They are also able to form triple helices in which a first CNA strand binds with RNA or ssDNA and a second PNA strand binds with the creater soluble, which facilitates cellular cuptake. Further, since they contain anddes of non-biological amino acide, they are biostable and resistant to enzymatic degradation by proteases. CC The present sequence is a specifically claimed PNA sequence (represented CC by the sequence of nucleobases) targetting the HIV gag/pol frameshift. Oligomer hybridisable to HIV sequence and contg. peptide nucleic acid sub:unit - binds in complementary manner to DNA and RNA, and useful for modulating HIV viral activity, e.g. in treating AIDS. WPI; 1995-082179/11 Claim 2; Page 177; 186pp; English. (ISIS-) ISIS PHARM INC 93US-00099718 94WO-US008517

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Sequence
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A; 5
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4 T; 0
U; 0 Other;
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RESULT 2980

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RESULT 2979
AAT41212
ID AAT4121
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Best Local S
                                                          Matches
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Best Local (
                                                                                                                                       Primers T41001-T41382 are derived from novel human gene signature (GS) sequences which did not match with sequences deposited in Genbank release 76. The GS sequences (T19001-T26837) were obtained from 3'-directed cDNA libraries prepared from various human tissues; synthesis of cDNA was initiated from the 3'-end of mRNA by using poly(T) as the sole primer. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was derived. The appearance frequency of a given GS in a cDNA library can be determined (esp. using primers and probes derived from the GS sequences) as means of diagnosing abnormal cell function or for recognising different cell types. The primers T41211-2 amplify clone pm0647 which comprises the GS HUMGS001132 (T20132), located on chromosome 20
                                                                                                                                                                                                                                                                                                                                                        Single-stranded DNA for identifying gene signatures - isolated from 3'-directed human cDNA library that reflects relative abundance of corresp. _{\rm mRNA} in specific human tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene signature; messenger RNA; mRNA; relative abundance; frequency; human; cloning; mapping; non-biased library; diagnosis; detection; cell typing; abnormal cell function; primer; PCR; amplification;
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                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                 Example 7; Fig 8; 2245pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Matsubara K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MATS/) MATSUBARA K. (OKUB/) OKUBO K.
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                            1936 ATCCTAGTCCCACACCAG 1953
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                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     signature HUMGS01132-derived anti-sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chain reaction; ss.
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                                                                                                               BP; 6 A; 8 C; 2 G; 4 T; 0 U; 0 Other;
                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                 Japanese.
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                                                          Pred. No. 2e+
0; Mismatches
                                                                                   Score 14.8;
                                                                         No.
                                                                       2e+03;
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                                                                                   DB 1;
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RESULT 2981
AAT33418
ID AAT3341
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           translation termination. The second mutation is a G to A substitution at a splice donor site. The mechanism of how this mutation causes factor XIII deficiency is yet to be determined. The third mutation is a nonsense mutation in which a C to T transition at position 598, in an Arg codon, results in a stop codon TGA. A further eight mutations have been identified and include a deletion/insertion event, a nonsense mutation and missense/silent mutations. These primers may be used in the diagnosis and treatment of disorders involving factor XIII e.g. bleeding,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             products obtained for each exon of each individual were found to be of the expected size, indicating that there are no gross insertions or deletions in the facto XIII "a" gene of these patients. Three mutations which may be the cause of "a" subunit deficiency have been described. The first is a two base pair deletion at a splice donor acceptor site. This deletion does not grossly affect the splicing of the factor XIII pre mRNA, but causes a translational frameshift resulting in early
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequences given in AAT28903-32 are primers which were used in the amplification of the exons of the factor XIII "a" gene. This allows analysis of the factor XIII gene and identification of differences in gene sequence which are known to segregate with a reduction or enhancement of factor XIII activity. All fifteen exons were amplified from five unrelated families showing factor XIII disorders. The PCR from the contract of the factor XIII activity and factor XIII disorders.
   AAT33418 standard; cDNA; 20
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splice donor/acceptor site; translational frameshift; substituti
nonsense mutation; transition; diagnosis; bleeding; haemorrhage;
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genetic study of Factor XIII activity - used for diagnosis and treatment of Factor XIII disorders, e g bleeding, haemorrhage, miscarriage or clot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W09617953-A2
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16; Conserv
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                                                                                                                                                     TACTCCCAGCAACTGGTT
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                                                                                                                                                                                                                                                                           Conservative
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Pred. No. 2e+0
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                                                                                                                                                                                                                                                                                                     2e+03;
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RESULT 2982
AAT33421
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Matches
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                                                                                                 Synthetic.
                                                                                                                                                   Antisense; VEGF; vascular endothelial growth factor; hypoxia; neovascularisation; angiogenesis; metastasis; retinopathy; macular; degeneration; expression inhibitor; ss.
                                                                                                                                                                                                                                                               Human vascular
                                                                                                                                                                                                                                                                                                                                                                      AAT33421;
                                                                                                                                                                                                                                                                                                                                                                                                                           AAT33421 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  essentially consist of 2'-O-alkylated ribonucleotides. Inhibiting expression of VEGF is useful in the treatment of hypoxia induced neovascularisation and angiogenesis associated disease states,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT33371-T33431 are antisense oligonucleotides used to inhibit the expression of human vascular endothelial growth factor (VEGF). The synthetic oligonucleotides contain phosphorothioate linkages and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 14; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human vascular endothelial growth factor anti:sense oligo:nucleotide - inhibits the expression of VEGF, useful in treatment of hypoxia induced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-MAR-1995;
08-DEC-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neovascularisation and angiogenesis associated disease states.
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expression inhibitor; ss.
                                                                                                                                                                                                                                                            endothelial growth factor
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95US-00569926
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                                                                                                                                                                                                                                                                                                                                                                                                                           CDNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8;
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                                                                                                                                                                                                                                                               antisense
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06-SEP-1996

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RESULT 2983
AAT33412
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT33371-T33431 are antisense oligonucleotides used to inhibit the expression of human vascular endothelial growth factor (VEGF). The synthetic oligonucleotides contain phosphorothicate linkages and essentially consist of 2'-O-alkylated ribonucleotides. Inhibiting the expression of VEGF is useful in the treatment of hypoxia induced neovascularisation and angiogenesis associated disease states, retinopathy of prematurity, diabetic retinopathy and age related macular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human vascular endothelial growth factor anti:sense oligo:nucleotide - inhibits the expression of VEGF, useful in treatment of hypoxia induced neovascularisation and angiogenesis associated disease states.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-MAR-1995;
08-DEC-1995;
                                                                                                                                                                                                                                                                                                                                             16-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                           AAT33412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 14; 92pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-FEB-1996;
                                                                                                                                                                                                                                                                                                                   Human vascular endothelial growth factor antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
           neovascularisation
                      Human vascular endothelial growth factor anti:sense oligo:nucleotide -
inhibits the expression of VEGF, useful in treatment of hypoxia induced
                                                                                                                                02-MAR-1995;
08-DEC-1995;
                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                           Antisense; VEGF; vascular endothelial growth factor; hypoxia;
                                                                                                                                                                     29-FEB-1996;
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                                                                                                                                                                                                                     WO9627006-A2
                                                                                                                                                                                                                                                                    degeneration;
                                                                                                                                                                                                                                                                                 neovascularisation; angiogenesis; metastasis; retinopathy; macular;
                                                                                                         (HYBR-) HYBRIDON
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                                                          1996-412773/41
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                                                                                                                                                                                                                                                                                                                                                                                            standard; cDNA;
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                                                                                                                                                                                                                                                                    expression inhibitor; ss.
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95US-00569926.
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95US-00569926.
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            and
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          angiogenesis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 3 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8;
Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
            associated
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2e+03;
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            disease
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RESULT 2984
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Best Local (
                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT33371-T33431 are antisense oligonucleotides used to inhibit the expression of human vascular endothelial growth factor (VEGF). The synthetic oligonucleotides contain phosphorothicate linkages and essentially consist of 2'.O-alkylated ribonucleotides. Inhibiting the expression of VEGF is useful in the treatment of hypoxia induced neovascularisation and angiogenesis associated disease states, retinopathy of prematurity, diabetic retinopathy and age related macular
                                                                                                   Neovascularisation can be reduced by blocking vascular endothelial g factor (VEGF) expression using a synthetic oligonucleotide specific VEGF. Inhibiting neovascularisation is useful for treatment of retinopathy of prematurity, diabetic retinopathy and age-related mac degeneration. The present sequence is an example of a suitable oligonucleotide specific for human VEGF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT48412 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 14; 92pp; English.
                                                                                Sequence
                                                                                                                                                                                     Disclosure; Page 12; 66pp; English.
                                                                                                                                                                                                         Inhibiting neovascularisation using VEGF-specific oligo:nucleotide(s) for treatment of retinopathies and age-related macular degeneration.
                                                                                                                                                                                                                                            WPI; 1996-362689/36.
                                                                                                                                                                                                                                                                                         (HYBR-)
(CHIL-)
                                                                                                                                                                                                                                                                                                                            26-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                         01-AUG-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                             diabetes; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vascular endothelial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT48412;
                                                                                                                                                                                                                                                                   Robinson GS,
                                                                                                                                                                                                                                                                                                                                                  26-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        neovascularisation;
5921 CCCAGAGATGTCCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5921 CCCAGAGATGTCCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                   l Similarity
16; Conserv
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CHILDRENS MEDICAL
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                                                                               BP; 5
                                    Conservative
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                                                                                                                                                                                                                                                                    Smith
                                                                                                                                                                                                                                                                                                                            9505-00378860.
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                                            0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    growth factor; inhibition; decrease; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific
                                                                                9 C; 3 G; 3 T; 0 U;
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              5938
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Pred.
                                                         Score 14.8;
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                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        age-related macular degeneration;
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                                                                                0 Other;
                                                         DB 1;
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                                                         Length
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                                     Indels
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                                                                                                                              age-related macular
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                                     Gaps
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ARTSULT 2986
AAT48421
ID AAT4842
XX AAT4842
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XT 11-MAR-
XT 0ligonu
XX Vascula
KW vascula
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XX XX
OS Synthet
XX
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AAT48418
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                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                AAT48421 standard;
                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                      factor (VEGF) expression using a synthetic oligonucleotide specific for VEGF. Inhibiting neovascularisation is useful for treatment of retinopathy of prematurity, diabetic retinopathy and age-related macular degeneration. The present sequence is an example of a suitable oligonucleotide specific for human VEGF
                                 neovascularisation; retinopathy; age-related macular degeneration; diabetes; exon-intron boundary; ss.
                                                        Vascular endothelial growth factor; inhibition; decrease; antisense;
                                                                               Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-362689/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JAN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diabetes; exon-intron boundary; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor; inhibition; decrease; antisense; neovascularisation; retinopathy; age-related macular degeneration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide
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           Synthetic
                                                                                                      11-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                               Neovascularisation can be reduced by blocking vascular endothelial
                                                                                                                                                                                                                                                                                                                                                                                                                           Inhibiting neovascularisation using VEGF-specific oligo:nucleotide(s) for treatment of retinopathies and age-related macular degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robinson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JAN-1996;
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(CHIL-) CHILDRENS MEDICAL CENT.
                                                                                                                                                                                                                              2402 CTGGGACCACAGTGGACA 2419
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                                                                                                    (first entry)
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                                                                              E3/I3 specific
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                                                                                                                                                DNA;
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Pred. No. 2e
                                                                               for human
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                                                                              VEGF nucleic acid
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                                                                                                                                                                                                                                                                         Length 20;
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AAV85793
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Best Local :
          Todd JA, Hess JW, Caskey CT, Hey P, Kawaguchi Y, Merriman Phillips MS, Twells RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neovascularisation can be reduced by blocking vascular endothelial growth factor (VBGF) expression using a synthetic oligonucleotide specific for VBGF. Inhibiting neovascularisation useful for treatment of retinopathy of prematurity, diabetic retinopathy and age-related macular degeneration. The present sequence is an example of a suitable oligonucleotide specific for human VEGF
                                                                                                                                                                                                                                                   LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endo
insulin dependent diabetes mellitus; autoimmune disease;
glomerulonephritis; inflammation; viral infection; ostooporosis;
                                                      (WELL )
                                                                                        15-APR-1997;
05-JUN-1997;
                                                                                                                                                                      WO9846743-A1
                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                             hypercholesterolemia; Alzheimer's disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                 LRP5 exon
                                                                                                                                                                                                                                                                                                                                               AAV85793;
                                                                                                                                                                                                                                                                                                                                                                    AAV85793 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                          15-APR-1998;
                                                                                                                                                 22-OCT-1998.
                                                                                                                                                                                                                                                                                                                         10-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Inhibiting neovascularisation using VEGF-specific oligo:nucleotide(s) for treatment of retinopathies and age-related macular degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-JAN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
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                                                      WELLCOME TRUST LTD.
MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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97US-0048740P
                                                                                                                          98WO-GB001102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 C; 3 G; 3 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8;
Pred. No. 2
                      TR,
                     Cox RD, Gerhol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2e+03;
                                 Gerhold D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                         low density lipoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>ب</u>
                     Nakagawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                  Hammond
                                                                                                                                                                                                                                                                           endocytosis;
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RESULT 2988
AAV85871
ID AAV8587
AC AAV8587
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85587 to AAV85822 represent exon primers used for obtaining LRP5 CDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated LDL-receptor related protein - used to develop products for treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
                                                                                                                                   Hey P, F
Phillips
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune diaease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
New isolated LDL-receptor related protein - used to develop products treating, e.g. elevated triglyceride levels, diabetes, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LRP5 SNP primer 58-8 lf.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-FEB-1999
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                                                                                                                                                                                                                                                                                                                                                                                                             15-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                   WPI; 1998-594573/50
                                                                                                                                                                                                                                             (WELL )
                                                                                                                                                                                                                                                                                                                           15-APR-1997;
05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9846743-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-OCT-1998
                                                                                                                                           J JA, new-
P, Kawaguchi '
Twel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer;
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o be
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                                                                                                                                                                                                                                                MERCK
                                                                                                                                                            Hess JW, co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                             ξ,
                                                                                                                                        Twells
                                                                                                                                                                                                                                                                                                                           97US-0043553P.
97US-0048740P.
                                                                                                                                                                                                                                                                                                                                                                                                               98WO-GB001102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                               Caskey CT,
Merriman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8;
Pred. No. 2e
                                                                                                                                                            Cox RD, Gerhold D, Hammorr, Metzker ML, Nakagawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2
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                                                                                                                                                                                              Hammond H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
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RESULT 2989
AAV52002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85823 to AAV85900 represent SNB primers used for obtaining LRP5 CDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cyrckine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention can also be used for detection, diagnosis and drug screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV52002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV52002 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2075 GCCGATACTGTGCTACTG 2092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12; Page 111; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCCAAGACTGTGCTACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inflammation or Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.8
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Polymorphic marker; allele-specific; probe; hybridisation; plant; hybrid certification; progeny; back-cross; hybrid; ancestry; corn; 02-DEC-1996; 07-MAR-1997; Synthetic. progeny; back-cross; Zea mays genome 02-FEB-1999 Lemieux (AFFY-) AFFYMETRIX INC 01-DEC-1997; 11-JUN-1998. WO9824796-A1 Ē Landry BS, (first entry) 96US-0032069P. 97US-00813507. 97WO-US021782 reverse PCR Sapolsky primer #298. ŖJ, Murigneux ; amplification; PCR pr genetic contribution; ; 88. PCR primer;

AAV51705-V52008 are reverse PCR primers used to amplify fragments of the Zea mays genome in order to detect polymorphic markers. Such markers can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution o

to trace the progeny of proprietary

Ö.

an ancestral plant,

Brassica species allele-specific useful for plant breeding.

oligonucleotide probes

and

primers

WPI; 1998-333252/29

Example

1; Page

55; 65pp; English.

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RESULT 2990
AAV35084/c
ID AAV3508
XX AAV3508
XX AAV3508
XX AAV3508
XX ARTisen
XX P-glyco
KW P-glyco
KW P-glyco
KW P-glyco
KW Primer;
XX W WO98146
XX W D1-OCT-
XX W WO98146
XX W D1-OCT-
XX W WO98146
XX W WPI; 19
PT OLOWING
XX W WPI; 19
PT ARTI-8e
PT ARTI-
  Query Match
Best Local S
Matches 16
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Matches 16; Conserv
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                                                                                                                                                                               AAV35061-V35101 are primers which have a sequence complementary to the translation initiation or termination region of a nucleic acid encoding a P-glycoprotein associated with multiple drug resistance (MDR) and inhibite expression of the glycoprotein. These primers are composed of 8-30 covalently linked nucleotides and includes at least 1 of the following, a 2'-modification, a lipophilic group (IG) that improves cellular uptake, and at least 1 covalent link that is a phosphorothioate, phospho di- or tri-ester, methylphosphonate, methylene (methylinino), morpholino, polyamide, short chain alkyl or heteroatomic inter-sugar link, or cycloalkyl or heterocyclic inter-sugar link. The primers are used to modulate human MDR gene expression in cells and tissues, i.e. to improve chemotherapeutic treatment of an animal with hyper-proliferative disease, particularly cancer, to prevent development of MDR and to re-
                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anti-sense oligo:nucleotide(s) targetted to multiple drug resistance gene - are modified by lipophilic substituent, on sugar and-or with non-natural linkages, used to improve activity of anti-proliferative agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-240109/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P-glycoprotein; multiple drug resistance; MDR; cellular uptake; cancer; gene expression; chemotherapy; treatment; hyper-proliferative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             certification of a crossed plant with
                                                                                                                                                                      sensitise an animal that has
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 21; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
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  Local Similarity es 16; Conserv
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                                                                                                              BP; 2
     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                       0.2%;
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                                                                                                              10 C; 4 G; 4 T; 0 U;
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Score 14.8; DB 1;
Pred. No. 2e+03;
0; Mismatches 2;
                                                                                                                                                                   developed MDR to a chemotherapeutic
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1103 AGAGTGGACAGACTGTGG 1120

RESULT 2992 AAV22586

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AAV70043/c
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                                                                                                                     AAV70042 to AAV70052 represent antisense oligonucleotides which are specifically hybridisable with a region of a nucleic acid encoding rat cross protein. The antisense compound regulates the expression of the c-Fos protein. The present invention also describes antisense oligonucleotides which regulate the c-Jun protein. The antisense oligonucleotides are used for the diagnosis and treatment of diseases or disorders associated with Activating Protein 1 expression, of which c-Fos and c-Jun are subunits. The antisense oligonucleotides are used in compositions as c-Fos and/or c-Jun together with a carrier and a chemotherapeutic agent. They are used to regulate the expression of c-Fos or c-Jun in cells or tissues, preferably by inhibiting metastasis. They also regulate cell cycle expression and can be used to treat an animal with, or being prone to, a hyperproliferative disease
                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotides regulating Activating Protein hybridise with c-fos and c-jun mRNA, used for regulating cycle expression and hyperproliferative disease.
                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-609906/51.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rat; c-fos; c-jun; activating protein 1; AP-1; diagnosis; metastasis; antisense oligonucleotide; phosphorothioate; regulation; malignant tumour; cell cycle expression; hyperproliferative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rat c-Fos
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                         1284 CCAGACCTCGACCATGAT 1301
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                                                            Similarity
                                                                                                                                                                                                                                                                                                       9; Page 57; 120pp; English.
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                                                          0.2%;
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RESULT 2993
AAV26329/c
ID AAV26329;
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AC AAV26329;
XX O7-AUG-199
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                                                                                                                                                                                                                                                                                                                                                                   CC AAV22531-89 represent antisense oligonucleotides which are targeted against the mRNA of the R1 subunit sequence of ribonucleotide reductase. CC Aberrant expression of the R2 gene, which encodes the second subunit of the ribonucleotide reductase gene, can determine the malignant contracteristics of cells. Suppression of R2 and R1 gene expression was CC found to reduce transformed properties of tumour cells. The antisense collgonucleotides can be used for modulating tumour cell growth, or for CC inhibiting tumour cell proliferation. They can also be used for CC increasing the sensitivity of neoplastic cells to chemotherapeutic drugs CC especially to hydroxyures, methotrexate (MTX), and pALA). The antisense coligonucleotides may be used to treat proliferative disorders including CC enter, papillomas, sarcomas, methodromas, various other forms of CC cancer, papillomas, sarcomas, methodromas, various other forms of CC cancer, papillomas, arthroscolarosis, psortassis, polythemia, mastocytosis, autonimune diseases, anglogenesis, bacterial infections and viral confections (including HIV hepatitis, or herpes infections)
                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R1 subunit; ribonucleotide reductase; cell proliferation; tumour cell; antisense; growth; inhibition; sensitivity; hydroxyurea; chemotherapeutic drug; methotrexate; PALA; treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-AUG-1996;
07-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense oligonucleotide designed to target the R1 message
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                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense oligonucleotides to ribonucleotide reductase genes - used modulate tumour growth and inhibit tumour cell proliferation.
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 Prostate cancer; human; marker; diagnosis; treatment; RT-PCR primer; ss
                                                                 07-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-145609/13.
                                Human prostate cancer marker UC Band #201 identifying RT-PCR primer
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                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                              CTTTTTTTTTTTTTTTTTTTTT
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                                                                                                                                 standard;
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97US-0039959P.
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can act as human prostate cancer markers are provided in the specification. The specification also provides methods for identifying markers for human prostate cancer and for detection of prostate cancer cells. The markers can be identified by amplifying human prostate RNA to provide nucleic acid amplification products, separating the products and identifying those RNA that are differentially expressed between human prostate cancers versus normal or benign human prostate. Prostate cancer cells in a sample can be detected by detecting a nucleic acid in a sample, the nucleic acid being a prostate cancer marker. Primers and probes derived from this marker can be used for the detection of prostate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This primer is used in the relative quantitative RT-PCR to exami expression of the genes which is used for the identification of of human prostate cancer. Isolated nucleic acid segments shown i payr16881 to AAV16885, AAV16890 to AAV16903, AAV26351 and AAV26352 and AAV26353 and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer cells in a sample. Antibodies against the protein encommarker nucleic acid fragments, inhibitors of the protein and oligonucleotides antisense to the markers can be used in the prostate cancer. The antibodies can also be used for the diag
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                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human prostate cancer
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                                                                                        3414 CTTATTCCTCTCTGTCCA 3431
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    19
                                                                                                                                                                                                                                  Similarity
CATATTCCTCTTTGTCCA 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 121; 229pp;
                                                                                                                                                                                                                                                                                                                                                                             BP; 9 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96WO-US012516
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                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                  2e+03;
                                                                                                                                                                                                                                                                                         DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the treatment of
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Query Match Best Local Sim Matches 16;

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RESULT 2994
AAX26076/c
ID AAX26076 standard; DNA; 20

Prostate disease marker gene fragment amplifying RT-PCR primer. 20-MAY-1999

(first entry)

diagnostic; Prostate cancer; differentiation; ncer; benign prostatic hyperplasia; marker gene; tumour; tion; Reverse Transcription Polymerase Chain Reaction; progression; cancer; metastasis; RT-PCR; primer; ss.

Homo sapiens Synthetic

US5882864-A.

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AAX29179
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to methods for diagnosing prostate cancer or benigr prostatic hyperplasia cells in a biological sample. The method uses oligonucleotides specific for marker genes associated with tumour differentiation and progression in Reverse Transcription Polymerase Chair Reaction (RT-PCR) analysis. The methods are diagnostic techniques useful for detecting and monitoring the progression of benign prostatic hyperplasia and human prostate cancer (the most prevalent form of cancer and a major cause of death in males) prior to the tumor undergoing metastasis, therefore allowing the optimal method of treatment to be determined before the condition becomes life threatening
                                                                                                                                                                                                                                   Osteopontin; antisense; restenosis; coronary arterial tissue; CASMC; inflammation; coronary artery smooth muscle cell; angioplasty; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diagnosing prostate cancer and benign prostatic hyperplasia cells - using oligonucleotide probes specific for marker genes associated with tumor differentiation and progression in Reverse Transcription Polymerase Chain
                       WPI; 1999-190049/16.
                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                        Human osteopontin
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                                                                                            07-AUG-1997;
                                                                                                                   07-AUG-1998;
                                                                                                                                          18-FEB-1999
                                                                                                                                                                 WO9907844-A2
                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                               18-JUN-1999
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                                                                    (USSH ) US
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osteopontin antisense
                                                                                                                                                                                                                                                                                                                                                                                                                              3414
                                                                                                                                                                                                                         probe;
                                                                                                                                                                                                                                                                                                                                                                                                        19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     CTTATTCCTCTCTGTCCA
                                                                                                                                                                                                                                                                                                                                           standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               analysis.
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                                                                      DEPT
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                                                                                                                                                                                                                                                                                              (first entry)
                                                                                          97US-0054967P
                                                                                                                   98WO-US016569.
                                              Kundu
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                                                                      HEALTH & HUMAN
                                                                                                                                                                                                                                                                        (OPN)
                                                                                                                                                                                                                                                                                                                                           DNA;
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                                                                                                                                                                                                                                                                     specific probe hOPN-P1.
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sequences -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8; I
Pred. No. 2e+0
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hara
                                                                      SERVICES
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U;
useful to treat restenosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prostate cancer or benign
ple. The method uses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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RESULT 2996
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                        28-NOV-1997;
17-DEC-1997;
04-NOV-1998;
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AAZ04197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to antisense osteopontin oligonucleotide sequences which are complementary to at least a portion of the human osteopontin (OPN) cDNA sequence (AAX29191). The antisense sequences are used to prevent restenosis in tissue, particularly coronary arterial tissue, especially where the patient is undergoing angioplasty, particularly percutaneous trans-luminal coronary angioplasty or directional coronary atherectomy. They prevent secretion of osteopontin by monocytes and macrophages which infiltrate to sites of inflammation following surgery. Osteopontin probably causes restenosis by inducing coronary artery smooth muscle cells (CASMC) to migrate to, and proliferate at, angioplasty injury sites. The present sequence represents a probe specific for human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
           against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as
                                       PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAX36754-X37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also
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                                                                                                                                                                                                                                                                                                                                                                                                                             Chlamydia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer used to amplify an ORF of Chlamydia trachomatis
                                                                                                                                                                                                                                           (GEST ) GENSET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA;
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                                                                                                                  Page 1669; 1755pp;
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97FR-00016034.
98US-0107077P.
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Pred. No. 2e+0
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                                                                                                                    English.
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RESULT 2997
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Best Local Similarity
                                                                                                                                                    PCR primers AA201426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY3673-4-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, peridpymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-NOV-1997;
17-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genome sequence of Chlamydia trachomatis.
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                                                                                        Sequence
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y Match 0.2%;
Local Similarity 88.9%;
hes 16; Conservative
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97FR-00016034.
98US-0107077P.
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Pred. No. 26
Score 14.8; DB 1; Length 20; Pred. No. 2e+03; o; Mismatches 2; Indels
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RESULT 2999 AAZ04201/c ID AAZ0420 XX AC AAZ0420

AAZ04201 standard; DNA; 20

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RESULT 2998
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Best Local Similarity
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17-DEC-1997;
04-NOV-1998;
                                                                                                                                           PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY3673-437349) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, pendymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
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Chlamydia trachomatis.
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                                                                                                                                                                                                                                                                                                                                         Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                               Griffais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccine; eye disease; conventional trachoma; nonendemic trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ04690;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ04690 standard;
                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-371125/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09928475-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer used to amplify an ORF
                                                                                                                                                                                                                                                                                                             Disclosure; Page 1709; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3638 AGGAGGTAGATGGGGAAG 3655
                       4013 AAATGAGAAAAAAGAGAG
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 20
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 AGAGGAGAAAAAAGAGAG
                                                                                                         BP;
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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97FR-00016034.
98US-0107077P.
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                                                                0.2%;
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                                                                                                         C; 1
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                          4030
                                                     0,
                                                                                                           G; 11 T; 0 U;
                                                  Score 14.8; DB Pred. No. 2e+03; O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of Chlamydia trachomatis.
                                                                                DB 1; Length
                                                                                                           0 Other;
                                                     2;
                                                      Indels
                                                      0,
                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer;
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RESULT 3000
AAZO1618
ID AAZO161
XX AAZO161
XX AAZO161
XX O7-OCT-
XX O7-CCT-
XX VACCINE
KW VACCINE
KW PARATER
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                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY36734-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-NOV-1997;
17-DEC-1997;
04-NOV-1998;
 Synthetic
                                          paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                               bartholinitis;
                                                                                                                                      07-OCT-1999
                                                                                                                                                                                               AAZ01618
                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 1669; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR
                                                                           Vaccine;
                                                                                                         PCR primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genome sequence of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-371125/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer used to amplify an ORF
                                                                                                                                                                                                                                                                                                      2851 CCAAATCCAGAGGAAGCA 2868
                                                                                                                                                                                                                                                                          20
                                                                           eye disease;
                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        eye disease;
                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                              BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97FR-00015041.
97FR-00016034.
98US-0107077P.
                               pneumopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-IB001939.
                                                                                                                                                                                                                                                                                                                                                                                              Α,
                                                                                                       amplify an ORF
                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                             4 C; 6
                                                                           conventional trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       conventional trachoma;
                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                          w
                                                                                                                                                                                                                                                                                                                                   0;
                               venereal lymphogranulomatosis; ss
                                                                                                                                                                                                                                                                                                                                                                                             G; 8 T;
                                                                                                                                                                                                                                                                                                                                                  Score 14.8;
Pred. No. 2e
                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                    of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                             0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                  2e+03;
                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                           nonendemic trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nonendemic
                                                                                                                                                                                                                                                                                                                                   ب
                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       trachoma;
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                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 3001
AAZ03279
ID AAZ0327
XX AAZ0327
XX AAZ0327
XX VAccine
KW PARACTA
KW PARACTA
KW nongonc
KW barthol
XX nongonc
KW barthol
XX NO99284
XX PB WO99284
XX PF 27-NOV-
XX 28-NOV-
PR 17-DEC-
PR 04-NOV-
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
 28-NOV-1997;
17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-NOV-1997;
17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY3673-4-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as mongonococcal uretritis, penitaly in breast feeding infants, and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                 Synthetic.
Chlamydia trachomatis.
                                                                                                                                                                              Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepat nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                              07-OCT-1999
                                                                                                                                                                                                                                                                                                                                   AAZ03279 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-371125/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              พO9928475-A2
                                                      27-NOV-1998;
                                                                                 10-JUN-1999.
                                                                                                             WO9928475-A2
                                                                                                                                                                                                                                                   PCR primer used to amplify an ORF of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                         AAZ03279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genome sequence of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEST )
                                                                                                                                                                                                                                                                                                                                                                                                                                 2652 CCACCTGGTGGACAAGGA 2669
                                                                                                                                                                                                                                                                                                                                                                                                       \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENSET
                                                                                                                                                                                                                                                                                                                                                                                                       CCACATGTTGGACAAGGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 1457; 1755pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                              (first
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97FR-00016034.
98US-0107077P.
              97FR-00015041.
97FR-00016034.
                                                      98WO-IB001939
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                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
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                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB Pred. No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                         perihepatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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                                                                                                                                                                                            primer;
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RESULT 3002
AAX9487
AAX9487
AC AAX9487
XX AAX9487
AC AAX9487
XX Respir
XX GALam
XX GALam
XX GES
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAX36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Griffais R,
     AAX91991-X97517 represent PCR primers used to amplify open reading and other nucleic acid sequences from the genome of Chlamydia pneum
                                                                                                                                                                                                       WPI; 1999-357842/30
                                                                                                                                                                                                                                                           Griffais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX94878 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-371125/31
                                                                                        Page 1704; Disclosure;
                                                                                                                                                                                                                                                                                                                                                                       21-NOV-1997;
04-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chlamydophila
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                                                                                                                                                                                                                                                                                                                 (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5426 AAGAGAATCAGCTTTGGG
                                                                                                                                              sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                           ۳,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGAGGATCAGCGTTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 1593; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pneumoniae
                                                                                                                                                                                                                                                                                                                                                                       97FR-00014673.
98US-0107078P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-IB001890
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 C; 8
                                                                                     1912pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8;
Pred. No. 2
                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
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These ORFs
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pneumoniae

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RESULT 3003
AAZ47569/c
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This sequence is an antisense oligonucleotide that specifically hybridises to nucleic acids encoding a human multidrug resistance P-glycoprotein (MDR1). The oligonucleotide inhibits expression of the P-glycoprotein, which functions as an ATP driven efflux pump. The antisen oligonucleotides of the invention have a phosphorothioate modified backbone, and may contain residues with 2' modifications selected from methoxyethoxy, 2'-fluoro, 2'-O-fluoro or 2'-propyl. Some antisense oligonucleotides have cholesterol bound at the 3' end which ensures resistance to 3' exonucleases, enhances cellular uptake, and leaves the 5'terminus available for conjugation of additional functional groups. To oligonucleotides may be used in research, diagnosis or as therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (see AXX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinustitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584- AAY35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                                                                                                                                                                                                                    Antisense oligonucleotide specific for multidrug resistance P-
glycoprotein is useful for treating hyperproliferative diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 8 A; 5 C; 5
                                                                                                                                                                                                                                                                                                                 Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                              04-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                           30-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Multidrug resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense oligonucleotide 24 targeted to human MDR1 P-glycoprotein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ47569 standard;
                                                                                                                                                                                                                        disorders e.g. cancer.
                                                                                                                                                                                                                                                                                   WPI; 2000-061907/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ug resistance gene; MDR1; human; hy
autoradiography; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                           Col 13; 24pp; English.
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                                                                                                                                                                                                                                                                                                                                                PHARM INC
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                                                                                                                                                                                                                                                                                                                 Dean NM;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hyperproliferative disease;
te; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
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                                      leaves the
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RESULT 3004
AAC69251/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-MAR-1999;
08-JUN-1999;
The invention relates to the human ABC1 cholesterol transporter protein (B38082) and to nucleic acid sequences (C69120) which encode it. ABC1 is a member of the ATP-binding cassette (ABC transporter) superfamily of proteins, and plays a crucial role in cholesterol transport, particularly intracellular cholesterol trafficking in monocytes and fibroblasts, being
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATP-binding cassette; HDL deficiency disorder; high density liporangier disease; TD; familial HDL deficiency; FHA; polymorphism; cardiovascular disease; coronary artery disease; coronary rester
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene expression can be used to treat hyperproliferative diseases and disorders e.g. cancer, in conjunction with chemotherapeutic reagents to prevent or modulate the development of multidrug resistance during the treatment. The oligonucleotides can also be used to resensitize hyperproliferative MDR cells in an animal previously exposed to chemotherapeutic agents. Radiolabelled oligonucleotides can be used to perform autoradiography of tissues to determine localization,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         distribution and quantitation of MDR P-glycoproteins diagnostic purposes
                                                                                                                                                                                                                                                                              WPI; 2000-587528/55
                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUN-1999;
01-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alzheimer's disease; Niemann-Pick disease; Huntington's disease;
X-linked adrenoleukodystrophy; cancer; gene therapy; genetic dia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human ABC1 cholesterol transporter; chromosome 9q31; ATP-binding cassette; HDL deficiency disorder; high
                                                                                                                                           Disclosure;
                                                                                                                                                                                                           New ABC1 po
biological
                                                                                                                                                                                                                                                                                                                           Hayden MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-SEP-2000
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                                                                                                                                                                                                         polypeptide is useful for treating diseases associated with all activity, e.g. Alzheimer's disease, Huntington's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGAGTGGACAGACTGTGG 1120
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                                                                                                                                                                                                                                                                                                                                                                   BRITISH COLUMBIA.
N BIORESEARCH INC.
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99US-0138048P.
99US-0139600P.
99US-0151977P.
                                                                                                                                       10; 229pp; English
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                                                                                                                                                                                                                                                                                                                           SN;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC disease, particularly coronary artery disease, but also cerebrovascular CC disease, coronary restenosis, and peripheral vascular disease. CC Conversely, a high level of HDL has protective effects against CC Cardiovascular disease. The invention provides genetic constructs and cardis and methods of gene therapy for the treatment or prevention of CC cardiovascular disease comprising the administration of an expression cC cardiovascular disease comprising the administration of an expression cC encompasses compounds which mimic ABC1 activity, compounds which CC stimulate ABC1 expression and methods of screening for such compounds. It further relates to methods for determining whether a patient has an CC abc1 gene. Human ABC1 proteins and nucleotides can be used to treat or prevent cardiovascular disease, especially coronary artery disease, cerebrovascular disease, especially coronary artery disease associated CC with ABC1 biological activity, such as Alzheimer's disease associated with ABC1 biological activity, such as Alzheimer's disease, and the exact sequence as GenBank Accession No: AJ012376.1. The invention specifically excludes proteins with the exact amino acid cand with the exact sequence as GenBank Accession No: AJ012376.1. The CC present sequence represents a human ABC1 gene PCR primer which may be used to amplify an exon of the human ABC1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            involved in cholesterol efflux from the cell. The gene encoding Abla Located on chromosome 9q31, and mutations in this gene are associated with two genetic HDL (high density lipoprotein) deficiency disorders, Tangjer disease (TD) and familial HDL deficiency (FHA). These diseases are distinguishable in that TD is an autosomal recessive disorder, while FHA is inherited as an autosomal dominant trait. Low levels of HDL ("good cholesterol") in the blood correlate with a high risk of cardiovascular cholesterol".
               US6046049-A
                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                    Phosphatidyl catalytic su
                                                                                              modified_base
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                                                                                                                                                                                                                                                                                                   diagnostic;
                                                                                                                                                                                                                                                                                                                                                                     PI3K antisense
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                                                                                                                                                                                                                                                                                                    lyl inositol 3 kinase; PI3K; antisense oligonucleotide; p110
subunit; treatment; rheumatoid arthritis; asthma; research;
; infection; inflammation; tumour formation; inhibitor; ss.
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                                                                                                                                                                                                                                                                                                                                                                     inhibitor oligonucleotide ISIS# 32136.
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1. .20
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                                                                base= OTHER
                                                                                                                "Optionally
                                                                                                                                                                                  "Phosphorothicate
                                               "Optionally 2'-methoxyethyl (2'-MOE)
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                                                                                                                                   OTHER
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Pred. No. 2
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                                                                                                                2'-methoxyethyl (2'-MOE) nucleotides"
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                                                                                                                                                                                  internucleoside linkage'
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                                                 nucleotides"
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RESULT 3006
AAA96416
ID AAA9641
XX AAA9641
XX O8-FEB-
XX O8-FEB-
XX AUCOIMM
KW ICOS ge
KW INSULIN
KW Graves
KW Craves
KW Craves
KW Craves
KW Craves
KW Craves
KW Chyroid
KW Chyroid
KW Chyroid
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XX Homo sa
XX WO20005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mitogenesis, protein trafficking, cell survival and proliferation, and chany other cellular activities. PIJK is a heterodimer, consisting of a continuous catalytic subunit (pll), and an 85kD regulatory subunit (pll), and an 85kD regulatory subunit (pll). The continuous catalytic subunit (pll), and an 85kD regulatory subunit (pll). The continuous catalytic subunit (pll) and the continuous catalytic subunit (pll) and the continuous catalytic subunit (pll). The continuous catalytic subunit (pll) and catalytic subunitation of catalytic subunitation (pll) and catalytic subunitation (pll) and catalytic subunitation of catalytic subunitation (pll) and catalytic subunita
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compounds targeting nucleic acids encoding human PI3 kinase p110 delta useful for treating a disease or condition associated with PI3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-APR-2000.
        WO200056856-A2
                                                     Homo sapiens.
                                                                                                                                                     Autoimmune disease; polymorphic microsatellite repeat; PMR; CD28 gene; ICOS gene; CTLA4 gene; costimulatory receptor gene locus; CGRL; lupus; insulin-dependent diabetes mellitus; IDDM; Addison's disease; leprosy; Graves disease; autoimmune hypothyroidism; myasthenia gravis; thymoma;
                                                                                                                                                                                                                                                                                                                                          08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA96416 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 16; Col 41; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-282691/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUL-1999;
                                                                                                                                                                                                                                                                                     Primer used
                                                                                                        thyroiditis; postpartum thyroiditis; rheumatoid arthritis; Hashimoto's disease; coellac disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3378 GTTGCTCCTCCCCAGCT 3395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   p110 delta expression, e.g. rheumatoid arthritis, asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                        to amplify a sara47/48 polymorphic microsatellite repeat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-00357070
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Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2e+03;
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3 kinases act
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RESULT 3007
AAZ87562/c
ID AAZ8756
AC AAZ8756
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC PCR primers AAA96415-16 were used to amplify polymorphic microsatellite crepeat (PMR) sequences from the human costimulatory receptor gene locus (hCGRI). The primers are used in the method of the invention. The specification describes a method for determining the predisposition of a pure subject to develop autoimmune disease. The method comprises contexting a pure sequence in the CD28, ICOS gene or CTIA4 gene of the chuman costimulatory receptor gene locus (hCGRI). PMR sequences vary in congrishment of the products and can be amplified to generate products that congrished the products can then be detected by rapid and convenient high resolution processes. The method is useful for convenient high resolution processes. The method is useful for convenient high resolution processes. The method is useful for convenient disease, craves disease, autoimmune hypothyroidism, compasthenia gravis, thymoma, lupus, thyroiditis, postpartum thyroiditis, convenient agravis, thymoma, lupus, thyroiditis, postpartum thyroiditis, conveniences within hCRGL are useful as markers in a variety of assays convenience within hCRGL are useful as markers in a variety of assays convenience within hCRGL are useful as markers in a variety of assays convenience.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                benign prostatic hyper; diagnosis; PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid marker; biomarker; tumour; prostate cancer; bladder cancer; benign prostatic hyperplasia; BPH; breast cancer; human; immunodetection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 1 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ling V,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer specific for cancer biomarker UC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ87562 standard; DNA; 20
                                                                                                                             11-JUN-1999;
                                                                                                                                                                                               16-DEC-1999
                                                                                                                                                                                                                                                          WO9964631-A1
                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEMY)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                   99WO-US013151.
                                                                                                                                                                                                                                                                                                                                                                                                                 hyperplasia; BPH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 C; 1
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Pred. No. 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Band #201.
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12-JUN-1998;

(UROC-) UROCOR

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RESULT 3008
ADA1128
ID AAA1128
XX AAA1128
XX O8-NOV-
XX O8-NOV-
XX Transme
KW Transme
KW Calcium
KW mutatio
XX Homo sa
XX Homo sa
XX Homo sa
XX WO20002
YM WO20002
XX WO20002
XX I1-NOV-
XX I1-NOV-
PR 11-NOV-
PR 11-NOV-
PR 11-NOV-
PR 11-NOV-
XX (EIKE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention provides nucleic acid markers of prostate, breast and CC bladder cancer. The markers are indicators of malignant transformation of CC prostate, breast and bladder tissues and are diagnostic of the potential CC for metastatic spread of malignant prostate tumours. The nucleic acid can also be used as targets for therapeutic intervention in prostate cancer. CC banign prostatic hyperplasia (BPH), bladder cancer or breast cancer. The markers may be used to design specific probes and primers, for the rapid CC analysis of prostate, bladder or breast biopsy samples. The probes and cprimers may also be used for in situ hybridization or in situ PCR CC detection and diagnosis. They may also be used to identify and isolate CC full length gene sequences form various DNA libraries. Antibodies against the polypeptide products of the markers can be used to treat prostate CC cancer, bladder cancer or breast cancer. The encoded proteins may be used to detect antibodies. The proteins and antibodies can be used in concerts, and for clinical diagnosis of these cancers. The antibodies may also be used for radioimaging to quantify and localize the encoded proteins
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                 mutation; ss.
                                                                                                                                                                                                                                                                                                                            Transmembrane
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                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA11286 standard;
                                          12-NOV-1998;
                                                                      11-NOV-1999;
                                                                                                   25-MAY-2000
                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                  AAA11286;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2;
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              (EIKE ) EIKEN
                                                                                                                               WO200029571-A1
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                                                                                                                                                                                                                                                                                                                                                           TRPC7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bladder
                                                                                                                                                                                                                                                                                                                                                         gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                            protein; TRPC7; brain; transient receptor potential;
el function; human; gene therapy; periodic psychosis;
                KAGAKU KK
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                                          98JP-00321200
                                                                      99WO-JP006289
                                                                                                                                                                     11. .20
/*tag= b
                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                 number= 1
                                                                                                                                                                                                                                                                                                                                                         intron
                                                                                                                                                                                                             *tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer.
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                                                                                                                                                                                                                                                                                                                                                        2 junction.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prognosis and management of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8; DB 1;
2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20
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AAZ73463/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to the isolation of a nucleic acid (AAA11284) coding for a transmembrane protein TRPC7 (AAY92944) which is expressed in brain and is homologous to transient receptor potential (TRP) protein. This suggests that the TRPC7 protein may have a calcium channel function. The genomic sequence has been shown to contain 31 introns. This sequence represents an exon/intron junction from the genomic TRPC7 sequence. The DNA and protein can be used in the diagnosis and treatment of disorders associated with TRPC7, especially the screening, monitoring and treatment (by gene therapy) of periodic psychosis, which appears to be associated with mutations in the TRPC7 gene
                                                                                                                                                                                                                                                                                                                                                  Human genome, biallelic marker; high density disequilibrium map, genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                WPI; 2000-013267/01.
                                                                                                                                         Cohen D,
                                                                                                                                                                  (GEST )
                                                                                                                                                                                           21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                21-APR-1999;
                                                                                                                                                                                                                                                           28-OCT-1999.
                                                                                                                                                                                                                                                                                   WO9954500-A2
                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                                                                                       diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                 Human biallelic marker upstream amplification primer
                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ73463 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding transmembrane protein TRPC7 expressed homologous to transient receptor potential protein useful in treatment of associated diseases such as periodic psychosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimizu N, Nagamine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ73463;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3676 ACCTCCAGCCAGAAAGCC 3693
                                                                                                                                                                                                                                                                                                             sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                 GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7; Page 38; 77pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACCTTCAGCAAGAAAGCC
                                                                                                                                         Blumenfeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 7 A; 6 C; 3 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                           98US-0082614P
98US-0109732P
                                                                                                                                                                                                                                 99WO-IB000822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                         Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                         Chumakov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                   ID NO: 7819.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           brain
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AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification

Novel biallelic markers used

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construct

a high density disequilibrium

human

genome

Page 1898;

2745pp;

English.

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RESULT 3010
AAC73675/c
ID AAC7367
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Best Local Sim:
Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                              The present sequence is an oligonucleotide used for antisense modulation of interleukin-5 (II-5) signal transduction. Oligonucleotides were designed to target nucleic acids encoding II-5 and II-5 receptor-alpha. The antisense oligonucleotides may be used for the treatment of diseases associated with II-5 signal transduction, II-5 expression or II-5 receptor-alpha expression. Such diseases include asthma and eosinophilic syndrome. The oligonucleotides are also useful for research uses and to prevent or delay infection, inflammation or tumour formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse; interleukin-5; IL-5; signal transduction; antisense oligonucleotide; antiasthmatic; immuno: IL-5 receptor-alpha; asthma; eosinophilic syndron
Sequence
                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide compound used to treat asthma and eosinophilic syndrome in humans modulates interleukin-5 signal transduction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAR-2000; 2000WO-US007318.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention
                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-594648/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200058512-A1
                                                                                                                                                                                                                                                                                          Example 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MY,
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  20 BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense
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                                                                                                                                                                                                                                                                                          Page 53; 156pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
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  ð,
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  9 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide
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  G; 5 T; 0 U;
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  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunosuppressive; cytostatic;
syndrome; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       #17981
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Query Match

0.2%;

Score 14.8;

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Length 20;

AAA90815 ID AAAS XX

AAA90815 standard; DNA;

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RESULT

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RESULT 3011
AAC60575
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                                                        Matches
                                                                       Best
                                                                                   Query Match
                                                                                                                                       The present sequence is one of a large number of antisense oligonucleotides which are targeted to nucleic acids encoding fra-1. The sequences may be oligodeoxyribonucleotides or chimeric oligonucleotides containing a central gap region consisting of ten 2'-deoxynucleotides, which is flanked on both sides by 2'-methoxyethyl (2'-MOE) wings. The oligonucleotides have a phosphorothioate backbone and the cytidine residues in the 2'-MOE wings are 5-methylsytidines. The fra-1 antisense oligonucleotides are useful for inhibiting the expression of fra-1 in human cells or tissues. They can be used for diagnostics, therapeutics, prophylaxis and as research respects and in kits. Use of the antisense prophylaxis and as research respects and in kits. Use of the antisense compounds may also be useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation
                                                                                                                                                                                                                                                                                                                                                                Novel antisense compound 8-30 nucleobases in length targeted to human -1 and which specifically hybridizes with and inhibits the expression human fra-1, useful for modulating the expression of fra-1 in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; fra-1; antisense oligonucleotide; phosphorothioate; cytostatic; antiinflammatory; 2'-methoxyethyl wing; 2'-MOE wing; infection; cancer;
                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                      Example 15; Col 42; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Taylor JK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
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                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-601552/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1999;
                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5272 ATAGGGAGCAGGTGGCAG 5289
                           4175 TAGGGAGGGGGTGGTTAT 4192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fra-1 mRNA antisense oligonucleotide ISIS 109066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 AGACGGAGCAGGTGGCAG
                                                        16;
                                                                       Similarity
                                                                                                                  20
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                                                                                                                  BP; 4 A; 1 C; 10 G; 5 T; 0 U; 0 Other;
                                                          Conservative
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                                                                      88.9%;
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                                                                                     0.2%;
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                                                                       Pred.
                                                          Mismatches
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                                                                         No.
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                                                                       2e+03;
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DOG; ge
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Canis f
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Best Local S
                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is an antisense oligonucleotide directed against the mRNA encoding the R1 component of mammalian ribonucleotide reductase. Ribonucleotide reductase catalyses the conversion of ribonucleotides to their corresponding decoxyribonucleotides and thus plays an important role in DNA synthesis and cell proliferation. Regulation of ribonucleotide reductase is altered in cultured malignant cells and increased levels of R2 protein and R2 mRNA have been found in pre-malignant and malignant tissues as compared to normal control tissue samples. The present antisense sequence is therefore useful for inhibiting tumourigenicity of neoplastic cells and inhibiting metastasis of tumour cells. It is also useful for increasing sensitivity of neoplastic cells to chemotherapeutic drugs, thus allowing chemotherapeutic treatments to be used in patients who have become resistant or less sensitive to chemotherapy. The sequence may be RNA or DNA and may comprise a modified backbone and/or nucleotide analonies.
                                               Dog; genome; genomic marker; radiation hybrid map; identification; chromosome location; gene marker; polymorphic microsatellite marker; phenotype; behaviour; pedigree; ss.
                Canis familiaris
                                                                                                                                                                                                            AAA66863 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotide; ribonucleotide reductase; R1 protein; R2 protein; tumour cell proliferation inhibition; cancer; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribonucleotide reductase R1 message antisense oligo AS-I-2769-20
                                                                                                            Dog genomic marker oligonucleotide sequence SEQ ID
                                                                                                                                               09-OCT-2000
                                                                                                                                                                              AAA66863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 32; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotide, AS-I-618-20, tumor cell growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-558216/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wright JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-FEB-2000; 2000WO-CA000120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENE-) GENESENSE TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA90815;
                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                            4463
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                                                                                                                                                                                                                                                                                                                              CTTTTTTTTTTTTTTT
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                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 0 A;
                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                           3 C; 1 G; 16 T; 0 U; 0 Other;
                                             gene marker;
pedigree; s
                                                                                                                                                                                                                                                                                                                          4480
                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                           Score 14.8; DB 1; Length 20; Pred. No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INC.
                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     is useful for inhibiting
                                                                                                                                                                                                                                                                                                                                                          2;
                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                             NO:725
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a radiation hybrid map of the dog (Canine familiaris) genome comprising the genome location of a marker selected from AAA66139 to AAA6642. The radiation hybrid map is useful for identifying and localising dog genes, since it covers approximately 80 % of the dog genome and provides a dense map integrating different types (i.e. Type I and Type II) of markers. The map and the dog genome markers (or complementary sequences) are especially useful to identify genes responsible for phenotypic and behavioural traits in dogs, to identify morbid genes, to analyse diseases and identify implicated genes in such diseases and their alleles, and to study dog pedigrees. They may also be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New radiation hybrid map of the dog, Canine familiaris, for e.g. identifying genes implicated in phenotypic and or in genetic diseases and for studying dog pedigrees.
                       Galibert F,
                                                                            13-NOV-1998;
                                                                                                     15-NOV-1999;
                                                                                                                                 25-MAY-2000.
                                                                                                                                                            WO200029615-A2
                                                                                                                                                                                     Canis familiaris.
                                                                                                                                                                                                            Dog; genome; genomic marker; radiation hybrid chromosome location; gene marker; polymorphic phenotype; behaviour; pedigree; ss.
                                                                                                                                                                                                                                                                    Dog genomic marker oligonucleotide sequence SEQ ID
                                                                                                                                                                                                                                                                                                09-OCT-2000
                                                                                                                                                                                                                                                                                                                                                    AAA66923 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 1 A; 8 C; 3 G; 8 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful for isolating corresponding human gene sequences e.g. genes involved in genetic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 84; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
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                                                  (CNRS ) CNRS CENT NAT RECH SCI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 Similarity
16; Conserv
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                                                                                                                                                                                                                                                                                               (first entry)
                       Andre C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CENT NAT RECH SCI
                                                                             98US-0108193P.
                                                                                                      99WO-IB001907
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Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                             map; identifica
microsatellite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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                                                                                                                                                                                                                                           identification;
                                                                                                                                                                                                                                                                    NO:785.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           behavioral traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dog (Canine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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WPI; 2000-387821/33

New radiation hybrid map of the dog, Canine familiaris, for e.g. identifying genes implicated in phenotypic and or in genetic diseases and for studying dog pedigrees.

genome, useful behavioral traits

Claim

87;

English.

Sequence 20

B₽; genetic

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8 C; 0

<u>ი</u>

7 T; 0 U; 0 Other;

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GGCTGGGGGTGGGAGGGG

18

involved

ä

diseases

The present invention describes a radiation hybrid map of the dog (Canine familiaris) genome comprising the genome location of a marker selected from AAA66139 to AAA66942. The radiation hybrid map is useful for identifying and localising dog genes, since it covers approximately 80 % of the dog genome and provides a dense map integrating different types (i.e. Type I and Type II) of markers. The map and the dog genome markers (or complementary sequences) are especially useful to identify genes responsible for phenotypic and behavioural traits in dogs, to identify morbid genes, to analyse diseases and identify implicated genes in such diseases and their alleles, and to study dog pedigrees. They may also be useful for isolating corresponding human gene sequences e.g. genes involved in genetic diseases

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RRESULT 3
ADAS1212
ID ADASX
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Best Local S
Matches 16
                                                                                                                                                                                                                                       Composition for treating hormone-regulated cancer, e.g. breast and prostatic tumors, comprising an antisense oligonucleotide that inhibits expression of insulin like growth factor binding protein-5 by hormone-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Insulin-like growth factor binding protein-5; IGFBP-5; human;
antisense oligonucleotide; hormone-regulated cancer; prostatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  UNIV BRITISH COLUMBIA. MIYAKE H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                  tumor cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IGFBP-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapy; ss.
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                                                                                                                                                    English
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Pred. No. 2
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2e+03;
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cc which inhibits expression of IGFBP-5 by hormone-regulated tumour cells.

Cc The compositions is useful for delaying progression of hormone-regulated tumour cells such as prostatic cancer cells or breast cancer cells, to an androgen-independent state, by treating hormone sensitive tumour cells.

Cc with the antisense sequence which inhibits expression of IGFBP-5 by the tumour cells. The composition can also be used for treating a hormone-cc responsive cancer in an individual, and administering the composition to the individual after initiation of hormone-withdrawal to induce apoptotic cell death of hormone-responsive tumour cells, and therefore delaying the progression of hormone-responsive cancer cells, and therefore delaying the composition of also be used for inhibiting or delaying the metastatic boney progression of an IGF-1 sensitive tumour in a mammal, by administering the composition to inhibit the expression of IGFBP-5 by the hormone-responsive cancer cells, and therefore inhibiting or delaying composition to inhibit the expression of IGFBP-5 by the hormone-responsive cancer cells, and therefore inhibiting or delaying composition to inhibit the expression of IGFBP-5 by the hormone-responsive cancer cells, and therefore inhibiting or delaying composition to inhibit the expression of IGFBP-5 by the hormone-responsive cancer cells, and therefore inhibiting or delaying composition of the tumour 8888888888888888888888888888888

Sequence 20 BP; 1 A; 1 C; 15 G; 3 T; 0 U; 0 Other;

Matches Best Query Match Local 3623 GGGTGGGGGTGGGAGAGG 16; Similarity Conservative 0.2%; 3640 0; Score Pred. Mismatches 14.8; No. 2e+03; DB 1; <u>ب</u> Length Indels 20, <u>,,</u> Gaps 0

RESULT 3016
AAD06582/C
ID AAD0658
AC Human a
XX
Human;
KW pharmac
AC FIBR-)
AC 10-AUG-2001 AAD06582 standard; (first entry) DNA; ΒP

pharmaceutical; food industry; cosmetic; autoimmune disorder; vaccine; medical; arterial sealant; bone graft; dermal implant; haemostat; cancer; rheumatoid arthritis; beverage; photographic application; PCR primer; ss. Human; alphal(I) collagen; gelatin; cytostatic; viral alpha1(I) collagen gene coding region amplifying disorder; infection; SSCP 2REV primer.

sapiens.

WO200134647-A2.

17-MAY-2001.

10-NOV-2000; 2000WO-US030792

12-NOV-1999; 99US-00439058 10-NOV-2000; 2000US-00709700

(FIBR-) FIBROGEN INC

Bell MP, Neff TB, Polarek ÄĽ, Seeley

WPI; 2001-335911/35.

Novel isolated and purified bovine useful in medical, pharmaceutical, useful in vaccine, and for treating autoimmune disorders, infections or porcine collagens and gelatins food and cosmetic industries, as and cancer.

Example 1; Page 56; 168pp; English.

The present sequence is a PCR primer used for amplifying the coding region of human alphal(I) collagen gene. The present invention relate recombinant synthesis of collagens and gelatins derived from animals. Collagen is useful in medical, pharmaceutical, food and cosmetic industries. Collagen is an important component of arterial sealants, grafts, drug delivery system, dermal implants, haemostats, and incontinence implants, and for tracking autoimmune disorders such as rheumatoid arthritis. Collagen is useful in food products such as sau coding on relates sausage ö

This sequence represents an antisense oligonucleotide targeted agains human insulin-like growth factor binding protein-5 (IGFBP-5). The invention relates to a composition for treatment of hormone-regulated cancer, comprising an antisense oligonucleotide (such as this sequence

againet

45pp;

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AAK95028/c
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Best Local Similarity
Matches 16; Conser
                                                                                                                                                                                                                                                                                                                                                             08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 casings, and in cosmetics or facial and skin products such as moisturisers. Recombinant gelatin is useful in vaccine formulations for treating viral infections, autoimmune diseases and cancer. Gelatin is useful in the manufacture or as a component of various pharmaceutical an medical devices and products, in food and beverage industries, in hair care and skin care products, as a glue or adhesive in various manufacturing processes, as a light-sensitive coating in various electronic devices, as photoresist base in photolithographic processes, in printing and photographic applications, in laboratory application, an as a component in various gels used for biochemical and electrophoretic
                     clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA clone provided in the invention
                                                                                                                                                                                                                                                          WPI; 2001-524255/58
                                                                                                                                                                                                                                                                                       Wakamatsu
                                                                                                                                                                                                                                                                                                      Ota
                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUL-2000; 2000EP-00114089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; full length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human cDNA clone-specific primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-NOV-2001
                                                                                                                                                      The invention relates to primers for synthesising full length cDNA
                                                                                                                                                                                  Example 18; Page 129; 1380pp + Sequence Listing; English.
                                                                                                                                                                                                                                 830 Primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP1130094-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAK95028;
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                                                                                                                                                                                                                                                                                                                                  (HELI-)
                                                                                                                                                                                                                genetic
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su A, Sugiyama T, Nagai
                                                                                                                                                                                                                                                                                                                                 HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
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                                                                                                                                                                                                               manipulation.
                                                                                                                                                                                                                           useful for synthesizing
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2000JP-00118774.
2000JP-00183765.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA; cDNA synthesis; oligo-capping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 C; 6 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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Pred. No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                    Hayashi K,
K, Kojima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO: 4273.
                                                                                                                                                                                                                            full length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length
                                                                                                                                                                                                                                                                                     ŝ
                                                                                                                                                                                                                                                                                    Ishii S,
S, Otsuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
                                                                                                                                                                                                                              cDNA clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                    Kawai Y;
T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20;
                                                                                                                                                                                                                            and their use
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Query Match

0 .24;

Score 14.8;

DB 1;

Length

20;

15-NOV-2001 AAD15632;

(first entry)

RESULT 3019 AAD15632/c

AAD15632 standard;

DNA; 20

Sequence

20

ВP;

8

A; 7 C; 1 G; 4 T; 0 U; 0 Other;

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AAF87023
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 16; Conserva
                                         Matches
                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . .
                                                                                                                This sequence is a sequencing primer for the human CP2/LSF/LBP-1 gene ARNm. The invention relates to a method for predicting an increased risk of a human subject of developing Alzheimer's disease, comprising assaying for a mutation within the ADN sequence of the CP2/LSF/LBP-1 gene including the region controlling the expression of the gene. The method is useful for predicting an increased risk of a human subject of developing Alzheimer's disease. Transgenic animals containing sequences from the CP2/LSF/LBP-1 gene are useful for screening for drugs capable of
                                                                                   Sequence 20 BP; 4 A;
                                                                                                                                                                                                                                        CP2/LSF/LBP-1 gene.
                                                                                                                                                                                                                                                 Predicting increased risk of human developing Alzheimer's di
                                                                                                                                                                                                                                                                                                       Chartier-Harlin M,
                                                                                                                                                                                                                                                                                                                             (INSP )
                                                                                                                                                                                                                                                                                                                                                                                                                             EP1113081-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing primer
                                                                                                                                                                                                                   Example 1; Page 10; 35pp;
                                                                                                                                                                                                                                                                                    WPI; 2001-427121/46
                                                                                                                                                                                                                                                                                                                                                             28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                   28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                       04-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CP2/LSF/LBP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF87023;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF87023 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LBP-1;
                                                    Local
                   3140 ACTCTGTAGCCCTGCAGA 3157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                        l Similarity
16; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human;
                                                                                                                                                                                                                                                                                                                              INSERM
                                                                                                                                                                                                                                                                                                                                        INST PASTEUR LILLE
                                                                                                         or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AATGAGTTCATGTGTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTGAGTTCATGTGTGAG
                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene; sequencing primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  intron; Alzheimer's disease; diagnosis; ADN sequence;
                                                                                                                                                                                                                                                                                                                              INST NAT SANTE
                                                                                                                                                                                                                                                                                                                                                             99EP-00403304
                                                                                                                                                                                                                                                                                                                                                                                   99EP-00403304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for Human
                                                                                                                                                                                                                                                                                                        Amouye1
                                                                                  5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                         symptoms associated
                                                  88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.9%;
                                                              0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                   English.
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18
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                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CP2/LSF/LBP-1 ARNm sequence.
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                                                     Pred.
                                                              Score 14.8;
                                                                                                                                                                                                                                                                                                                              & RECH MEDICALE.
                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                     No.
                                                  2e+03;
                                                                                                         with Alzheimer's
                                                              DB 1;
                                        2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2;
                                                             Length
                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                               20;
                                                                                                                                                                                                                                                             disease,
                                                                                                         disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                        Gaps
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RESULT 3020
AAH28626
ID AAH2862
XX AAH2862
XX AAH2862
XX Human i
XX Human;
KW Himan;
KW inflamm
KW fibrosi
KW 88.
OS Homo 88
XX W020012
XX W020012
XX W020012
XX PR 27-SEP-
XX 28-SEP-
                                                                                                                                                                                                                                                                                                                                                                                                             문
                                                                                                                                                                                                                                                                                                                                     S
                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method for selecting optimal subsequence antisense targets. The method involves preparing an antisense oligonucleotide capable of inhibiting mRNA expression of target mRNA sequences, as well as antisense oligonucleotides capable of binding DNA. The antisense and antisense libraries are useful for preparing therapeutic agents for the treatment of genetic disease. The present DNA sequence is human Bcl-2 protein target DNA related to the invention. Note: The present sequence is shown as DNA in the specification; however, in vivo,
                                                                                                                              Human; interleukin-13; IL13; single nucleotide polymorphism; SNP; cance inflammation; immune disorder; cytokine; asthma; chromosome 5q31; fibrosis; forensic; disease susceptibility; drug screening; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Selecting
 28-SEP-1999;
                       27-SEP-2000; 2000WO-US026556
                                                05-APR-2001
                                                                                                                                                                                                        17-JUL-2001
                                                                                                                                                                                                                                                      AAH28626 standard;
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-529916/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BOLL/)
(GRAY/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-FEB-2000; 2000US-00504653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-AUG-2001
                                                                      WO200123410-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200161030-A2
                                                                                             sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                    target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bc1-2
                                                                                                                                                                               interleukin-13 coding
                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                       40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AP,
                                                                                                                                                                                                                                                                                                                                                              l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GRAY D M.
JU-SEOG L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bc1-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9; Page 28; 87pp; English
                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                AGGCCCCGCGCGCGCCC
                                                                                                                                                                                                                                                                                                                                       AGGCTCCGCGGCGGCGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         optimal subsequence n of target mRNA for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gray DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein target
                                                                                                                                                                                                                                                                                                                                                                                                           BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence would be mRNA
                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001WO-US004732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein;
                                                                                                                                                                                                        (first entry)
 99US-0156489P
                                                                                                                                                                                                                                                                                                                                                                                                            ₽,
                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                        0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                            10 C; 9 G; 1 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genetic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ju-Seog
                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA
                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                w
                                                                                                                                                                                                                                                                                                                                       57
                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                sequence fragment PCR primer
                                                                                                                                                                                                                                                                                                                                                                        Score 14.8;
Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense targets for inhibition of mRNA the therapeutic treatment of genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #6
                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense target; therapeutic;
                                                                                                                                                                                                                                                                                                                                                                        2e+03; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                #
                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                       cancer;
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S
                                  Query Match
Best Local S
Matches 16
                                                                                                                                              human interleukin-13 (III3), and describes the single nucleotide polymorphisms (SNPs) found within the gene, which is found on chromoson 5q31. II13 is a pro-inflammatory cytokine thought to be involved in the pathogenesis of asthma and other immune and inflammatory diseases. The III3 sequences and the SNPs identified can be used in drug screening, t determine an individual's susceptibility to disease, in forensic and paternity testing, and to identify treatments for cancer, immune and inflammatory diseases, including asthma and diseases characterised by
                                                                                                                                                                                                                                                                                                                                                 Novel polynucleotide comprising single nucleotide polymorphisms in human interleukin-13 gene is useful for studying expression and function of interleukin-13, as well as diagnosing and treating cancer, inflammatory, and immune disorders.
                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                  The present invention provides the protein,
                                                                                                                                                                                                                                                                                                                  Example 1; Page 30; 85pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-343160/36
                                                                                                                                  fibrosis. The present sequence is an IL13 fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENA-)
     6866 CCTGGGCAGGGAGAGAGG
                                    16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENAISSANCE PHARM
                                                   Similarity
                                     Conservative
                                                                                                 BP; 6 A; 4 C; 9 G; 1 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RR,
                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nandabalan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INC.
                                                                                                                                                                                                                                                                                                                    English
   6883
                                  0
                                                 Score 14.8; DB
Pred. No. 2e+03;
                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stephens JC;
                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                  cDNA and genomic sequences
                                                                  <u>ب</u>
                                     <u>ب</u>
                                                                  Length
                                                                                                                                  PCR primer
                                     0
                                     Gaps
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of.

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RESULT 3021 AAH28642 문 AAH28642 standard; DNA; 20 3 CCTGAGCAGGCAGAGAGG BP 20

Human 17-JUL-2001 AAH28642; interleukin-13 (first entry) coding sequence fragment

Human; interleukin-13; IL13; single nucleotide polymorphism; SNP; cancer; inflammation; immune disorder; cytokine; asthma; chromosome 5q31; fibrosis; forensic; disease susceptibility; drug screening; PCR p

PCR

primer

#17

Homo sapiens

WO200123410-A2

05-APR-2001.

27-SEP-2000; 2000WO-US026556.

28-SEP-1999; 99US-0156489P.

(GENA-) GENAISSANCE PHARM INC

Chew P Denton RR, Nandabalan ٣ Stephens

WPI; 2001-343160/36.

Novel polynucleotide comprising single nucleotide polymorphisms in human interleukin-13 gene is useful for studying expression and function of interleukin-13, as well as diagnosing and treating cancer, inflammatory, disorders. diagnosing

Example 1; Page 32; 85pp; English

S 멍

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Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides the protein, cDNA and genomic sequences on human interleukin-13 (IL13), and describes the single nucleotide polymorphisms (SNPs) found within the gene, which is found on chromosome 5q31. IL13 is a pro-inflammatory cytokine thought to be involved in the pathogenesis of asthma and other immune and inflammatory diseases. The IL13 sequences and the SNPs identified can be used in drug screening, to sequences and the SNPs identified can be used in forensic and informatic and sequences and the SNPs identified can be used in forensic and individual's susceptibility to disease, in forensic and
The sequence represents nucleic acid biomarker UC band 201 primer #2, used in detection of prostate, breast and bladder cancer. Biomarker nucleic acid sequences can be used as hybridisation probes and primers that specifically hybridise to prostate cancer, benign prostatic hyperplasia (BPH), bladder cancer or breast cancer markers. Proteins encoded by the nucleic acid markers can be used to produce antibodies for the detection of prostate, breast or bladder cancer. The nucleic acids can be used as targets for therapeutic intervention in these diseases, in the identification and isolation of full-length gene sequences, including regulatory elements for gene expression, from genomic human DNA libraries, as hybridisation probes for screening genomic chuman DNA libraries. The kits comprising the nucleic acid sequences are useful for detecting bladder, breast or prostate cancer cells in a biological sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-JUL-1995;
11-JAN-1996;
31-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                              and bladder cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate; breast; benign prostatic |
                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acids as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biomarker UC band 201 primer #2 used in diagnosis/prognosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS03979;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US6218529-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS03979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fibrosis. The present sequence is an IL13 fragment PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UROC-) UROCOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    O'hara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                                                      4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTGGGCAGGGAGAGGG 6883
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCTGAGCAGGCAGAGAGG
                                                                                                                                                                                                                                                                                                                                 Col 71;
                                                                                                                                                                                                                                                                                                                                                                                                      prognosing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-0001655P.
96US-0013611P.
96US-00692787.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INC
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hyperplasia; BPH; therapeutic; humar
                                                                                                                                                                                                                                                                                                                            78pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ralph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                  biomarkers and targets useful for detecting, ng, and in developing treatments for prostate, breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.2%;
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                                                                                                                                                                                                                                                                                                                                 English.
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Pred.
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No. 2e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8; DB 1;
2e+03;
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Matches
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                                                                                                                                                                                                                    29-MAY-1999;
29-MAY-1999;
29-MAY-1999;
07-APR-2000;
                                                                                                                                                                                                                                                                                                                              ITS; internal transcribed spacer region; Mycobacterium fortuitum; Mycobacterium chelonae; Mycobacterium abscessus; Mycobacterium va. Mycobacterium flavescens; Mycobacterium asiaticum; tuberculosis; Mycobacterium porcinum; Mycobacterium acanilania.
                                                                                                                                                                                                (SJHI-)
                                                                                                                                                                                                                                                29-MAY-1999;
29-MAY-1999;
                                                                                                                                                                                                                                                                                                            Mycobacterium malmoense.
                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                WO200073436-A1
                                                                                                                                                                                                                                                                                                                           Mycobacterium
                                                                                                                                                                                                                                                                                                                                                                                  19-MAR-2001
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                                                                                                                                                                                          (PARK/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             3414 CTTATTCCTCTCTCTCA 3431
                                                                                                                                                              2001-061527/07
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                                                                                                                                                                                        SJ HIGHTECH
KIM C M.
PARK H K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                           Park HK,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                CATATTCCTCTTTGTCCA
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                                                                                                                                                                                                                                                                     2000WO-KR000477
                                                                                                                                                                                                                                                                                                                          porcinum; Mycobacterium acapulcensis; identification; diernhoferi; PCR primer; probe; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                  (first
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99KR-00019632.
99KR-00019633.
99KR-00019634.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.8; DB:
Pred. No. 2e+03;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                    of Mycobacterium
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                                                                                                                                                                                                                                                                                                                                                                    malmoense.
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Novel oligonucleotide sequences of internal transcribing spacer region non-tuberculosis mycobacteria (NTM) used as probes or primers for detecting and identifying mycobacteria and distinguish TB complex from of f

27; Page 65; 89pp; English

The present sequence is an oligonucleotide developed using a CC Mycobacterium ITS (internal transcribed spacer region) nucleotide CC sequence. ITS DNA sequences from M. fortuitum, M. chelonae, M. abscessus, CC M. vaccae, M. flavescens, M. asiaticum, M. porcinum, M. acapulcensis, M. CC diernhoferi genes were identified. The oligonucleotides derived from CC these sequences were used to develop PCR primers and hybridisation probes for detection and identification of Mycobacterium. ITS has a more CC polymorphic region than 16S rRNA and also has a conserved region. It is therefore highly effective as a target DNA for distinction of genotype. CC therefore highly effective as a traced to solid substrate, hybridise only considered indentify the specific mycobacteria, and thus they CC with nucleotide sequences in ITS of specific mycobacteria, and thus they can detect and identify the specific mycobacteria sensitively. The CC can detect and identify the specific mycobacteria mycobacteria coligonucleotide can also detect and identify the specific mycobacteria. CC by PCR amplification. Using the oligonucleotide primers or probes made CC from ITS of mycobacteria, it is possible to detect mycobacteria,

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RESULT 3024
AAF91363/c
ID AAF9136
XX AAF9136
XX AAF9136
XX AAF9136
XX Human E
XX Human E
XX Human E
XX Homo 88
XX US61875
XX Homo 88
XX US61875
XX Homo 88
RESULT 30
ABL57890
ID ABL5
XX
AC ABL5
XX
AC ABL5
XX
AC ABL5
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 distinguish tuberculosis (TB) complex from non-tuberculosis mycobacteria (NTM), and to identify mycobacteria species accurately and effectively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
  11-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to antisense compounds up to 30 nucleobases in length targeted to a E2F transcription factor 1 The invention is useful for inhibiting the expression of E2F transcription factor 1 in cells or tissues. The antisense oligonucleotides may also be used as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense compound capable of inhibiting the expression of E2F transcription factor 1, useful for preventing or delaying infection, inflammation or tumor formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-190981/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAR-2000; 2000US-00517584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human E2F transcription factor 1 antisense oligonucleotide #69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-MAY-2001
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                                                  ABL57890;
                                                                                                ABL57890
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 14 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
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                                                                                                standard;
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                                                                                                                                                                                                                        GGAGGAGGGGTGTGGGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     agent and to prevent infection, inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brown-Driver VL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Col 43; 40pp; English.
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  (revised)
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                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1;
Pred. No. 2e+03;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 3026
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                    example from field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to new Xanthomonas campestris bacteria strains made non-virulent by inactivation of at least one virulence gene but which have retained the capacity to produce exo-polysaccharides (preferably xanthan gum). One such virulence gene deleted to produce the bacterial strains was the hrpC2 gene (Hypersensitive Reaction and Pathogenicity). The hrp genes are essential for pathogenicity in plants. The present sequence is a PCR primer used to clone the hrpC2 gene in an example from the invention. (Updated on 11-SEP-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New Xanthomonas campestris bacteria strains for use in production of exopolysaccharides are made non-virulent by inactivation of at least one
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR; primer; hypersensitive exo-polysaccharide; xanthan
                                                                                                                         expression inhibition;
chronic obstructive pul
                                                                                                                                      Human CLCA1; goblet cell; mouse Gob-5 orthologue; drug screening;
expression inhibition; antisense therapy; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-JUN-2000; 2000WO-FR001725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Xanthomonas campestris; pv vesicatoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypersensitive reaction and pathogenicity, hrpC2, PCR primer XcC2.4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-JUL-2002
                                                                                                                                                                       Human CLCA1 sequencing primer PR22,
                                                                                                                                                                                                   11-SEP-2001
                                                                                                                                                                                                                          AAH46127;
                                                                                                                                                                                                                                                  AAH46127 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 25; 33pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        virulence gene.
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                                                                                    Homo
                                                                                                          sequencing primer; ss.
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                                                                                   sapiens.
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                                                                                                                                                                                                                                                                                                                                                              l Similarity
16; Conserv
                                                                                                                         obstructive pulmonary disease; bronchial asthma; antiasthmatic;
                                                                                                                                                                                                                                                                                                               TTCCACCTGGTCGACAAG
                                                                                                                                                                                                                                                                                                                                                                                                               20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                            Score 14.8
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                      14.8;
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                                                                                                                                                                          SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                          NO:29
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22-NOV-2000; 2000WO-JP008232.

31-MAY-2001.

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AAH46128
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Matches 16
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              New antisense nucleotide, useful for treatment and prevention bronchial asthma and chronic obstructive pulmonary disease.
                                                                                                                                                                                                                                                                       24-NOV-1999; 99JP-00333479.
27-APR-2000; 2000JP-00127589.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequencing primer;
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                                                                                                                                                         Nakanishi A,
                                                                                                                                                                                                                                                                                                                                                        22-NOV-2000; 2000WO-JP008232
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human CLCA1; goblet cell; mouse Gob-5 orthologue; drug screening; expression inhibition; antisense therapy; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH46128;
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                                                                                                        WPI; 2001-355935/37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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27-APR-2000;
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nes 16; Conserv
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2000JP-00127589.
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8; DB 1;
2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bronchial asthma; antiasthmatic;
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                                                 유
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Best Local S
Matches 16
The sequence represents primer c816F, used to sequence the human RAD51 gene. The nucleic acid is useful in diagnosing genetic predisposition or susceptibility to breast cancer in an individual using the following steps: (1) detecting a mutation in the RAD51 gene in a human subject, comprising analysing a sample from the subject to detect the mutation; (2) assessing the risk of developing breast cancer, comprising: (a) analysing a sample from the subject for the presence of BRCA1 and/or BRCA2 mutations; and (b) if (a) is positive, analysing the sample for a mutation in the RAD51 gene, where the presence of the RAD51 mutation indicates an increased risk in developing breast cancer in the subject compared to a subject having at least one of the BRCA mutations and a wild-type RAD51 gene. Primers derived from the sequence can be used in a kit for detecting a mutation in the RAD51 gene of a subject, which is associated with a predisposition to breast cancer, comprising at least 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gob-5 gene (coding sequence shown in AN446101) or its human counterpart, the CLCA1 gene (coding sequence shown in AN446102). The invention also relates to an antibody specific for the Gob-5 protein, medical and diagnostic compositions containing the antisense nucleotide or the antibody, and methods and kits for screening for compounds which inhibit the protein. Gob-5 and CLCA1 are proteins expressed by goblet cells. The antisense oligonucleotides and antibody are therefore useful for the treatment and prevention of bronchial asthma and chronic obstructive pulmonary disease. Sequences AN446135-AN446136 represent primers used in an exemplification of the invention to sequence human CLCA1 cDNA (AN446124)
                                                                                                                                                                                                                                                                                                                                                                                                           Wang
                                                                                                                                                                                                                                                                                                              New nucleic acids comprising a mutant of the RAD51 gene, useful for diagnosing genetic predisposition or susceptibility to breast cance
                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-235217/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-1999;
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                                                                                                                                                                                                                                                                             Example; Page 18; 42pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               SERVICES
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Best Local
                                                                                                              The present invention describes a method for predicting the potential of an oligonucleotide to hybridise to a (complementary) target nucleotide sequence, involving identifying a subset of oligonucleotides within the predetermined number of unique oligonucleotides based on the evaluation of the parameter. Oligonucleotides in the subset are identified that are clustered along a region of the nucleotide sequence that is hybridisable to the target nucleotide sequence. The present sequence is an oligonucleotide described in the exemplification of the invention. (Updated on 11-SEP-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid primers derived from the RAD51 gene sequence
                                                                                                                                                                                                                                                                               Predicting the potential of an oligonucleotide to hybridize to a target nucleotide sequence, useful for evaluating oligonucleotide probe sequences, by identifying a oligonucleotides based on the evaluation of
                                                                                                                                                                                                                                                                                                                                WPI; 2001-424456/45
                                                                                                                                                                                                                                                                                                                                                                                                    10-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                            10-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-SEP-2003
19-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH80771;
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                                                                                                                                                                                                                                                                      parameters
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                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                             (AGIL-) AGILENT TECHNOLOGIES INC.
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                                             l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                acid hybridisation; probe; primer; human; rabbit; HIV-1;
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                                                       0.2%;
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                                                                                          C; 1 G; 10 T; 0 U;
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Pred. No. 26
                                         Pred. No. 2e+0); Mismatches
                                                       Score 14.8; DB 1; Length 20; Pred. No. 2e+03;
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RESULT 3030

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6329 TGGGAACTTAGGCTTAAC

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Query Match Best Local S Matches 16

Similarity

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2e+03; DB 1;

Length 20 Indels

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Conservative

<u>,</u> Pred. Score 14.8; Mismatches

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The present invention relates to human and murine G protein-coupled CC receptor (GPCR) protein, IGPCR17 (ANMA8353 and ANMA8354). The coding CC sequence for IGPCR17 is useful in gene therapy for prevention, CC amelioration or treatment of diseases characterised by aberrant CC expression or activity of IGPCR17, where the disease is a psychiatric or CC central nervous system (CNS) disorder associated with signal processing in CNS such as learning and memory disorders, movement dysfunctions, CC tics, tremor, Tourette's syndrome, Parkinson's disease, Huntington's CC disease, dyskinesias, dystonia, pain and spasms. In addition, IGPCR17 and its coding sequence are useful in diagnosis, prevention, amelioration or treatment of diseases associated with signal processing in CNS, CC consessive compulsive disorder (CDD), multiple sclerosis, Alheimer's CC disease/dementia, anorexia, kidney diseases such as renal failure, CC disease/dementia, anorexia, kidney diseases such as renal failure, CC disease/dementia, anorexia, kidney diseases such as irritable bowel syndrome CC (IBS), diarrhoea, mortility disorders and conditions of delayed gastric comptying, osteoporosis, infections such as bacterial, fungal, protozoal cand viral infections, asthma, allergy, arthritis, sepsis and CC gynecological disorders. This sequence was used along with the primer of CC (IBS) for tissue-specific expression analysis of IGCR17 in an example of from the treatment of the presention manascale.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunosuppressive; gene therapy; psychiatric disorder; central nervous system disorder; movement dysfunction; schizophrenia; multiple sclerosis; Alzheimer's disease; kidney disease; obesity; gastrointestinal disorder; osteoporosis; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tranquiliser; antiparkinsonian; neuroprotective; nootropic; murine; anticonvulsant; metabolic; anorectic; anabolic; antiinflammatory; antidiarrheic; osteopathic; antiasthmatic; antiallergic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G protein-coupled receptor; IGPcR17; analgesic; neuroleptic; tranquillser; antiparkinsonian; neuroprotective; nootropic; murine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New human or mouse G protein-coupled receptor protein, IGPcR17, useful for diagnosis, prevention, amelioration or treatment of central nervous system disorders such as Tourette's syndrome, Parkinson's disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gynecological disorder; PCR primer; 88.
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Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7; Page 40; 71pp; English.
                                          invention. The resulting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wattler S,
    ВP;
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  4 A; 4 C; 6
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    G; 6 T; 0 U; 0 Other;
                                              PCR
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                                          product
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                                          is given
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ABK94887
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                                                                                                                                                                 are fat regulated. (I) or the polymucleotide encoding it (II) are useful for diagnosing the presence of or a predisposition for a disorder convolving fatty acid regulated genes in a subject. A composition containing (I) or (II) is useful for treating a disorder involving fatty acid (PUFA) disorder is selected from a polymusaturated fatty acid (PUFA) disorder, eczema, cardiovascular convolving fatty disorders (coronary artery disease, cerebrovascular disease or peripheral vascular consequence, including pancreatitis, estema, pancreatitis, estematorial solving fat research agent and diabetic complications. (I) or (II) is useful as research agent and materials for complications. (I) or (II) is useful as research agent and materials for constructing nucleotide probes and primers, expressed to isolate polymorphism, for detecting changes in the level of expression of (II), and as a diagnostic tool. This sequence represents a correct converse used to isolate polymorphism fat remains and disease.
                                                         Query Match
Best Local (
                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated polypeptide segment encoded by fat regulated genes, useful for diagnosing the presence of or a predisposition for a disorder involving fatty acid regulated genes in a subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sinusitis; asthma; pancreatitis; osteoarthritis; rheumatacne; body weight disorder; obesity; cachexia; anorexia; psychiatric disorder; cancer; cystic fibrosis; pre-menst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patty acid regulated gene; polyunsaturated fatty acid disorder; puph disorder; eczema; cardiovascular disorder; hypertriglyceridaemia; dyslipidaemia; atherosclerosis; coronary artery disease; cerebrovascular disease; peripheral vascular disease; inflammation;
                                                                                                                                                    for detecting genetic polymorphism, for detecting changes in the expression of (II), and as a diagnostic tool. This sequence reprepentation of the contract of the primer used to isolate DNA encoding fat regulated genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-NOV-2001; 2001WO-CA001632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diabetes; diabetic complication;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK94887;
                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (XENO-) XENON GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention describes an isolated polypeptide segment (I) whose genes
    3492 AGGCACTATGGCACTTAG 3509
                                      16;
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                                                           Similarity
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                                                                                                                    B₽;
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De Antueno
                                      Conservative
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                                                                                                                  5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pancreatitis; osteoarthritis; rheumatoid arthritis;
                                                         0.2%;
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Jenkins
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                                        Mismatches
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No. 26
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                                                         8; DB 1;
2e+03;
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Nwaka S
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polymorphism; PCR; primer; ss
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                                              Query Match
Best Local :
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                                                                                                                      (such as blood, pancreas or tongue) for sequence variations of the SAC1 gene. A sequence variation of the SAC1 locus may indicate a predisposition to diabetes, obesity and/or alcoholism and may provide a diagnostic mark. The polynucleotide can be detected in a biological sample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes
                                                                                                                                                                                                                                                                            The invention relates to an isolated polypeptide, comprising a variant form of mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obesity, diabetes or alcoholism associated with SAC1 expression. The sequences are useful in screening for drugs and sweeteners. Recombinant cell lines and transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; mouse; SAC1; carbohydrate; sweetener; ethanol; alcoholism; ss; obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas; blood; torgue; PCR primer; anorectic; antidiabetic; gene therapy; protein replacement therapy.
                                                                                                                                                                                                                               embryos may be used in screening for and identifying agents that induce or repress function of SAC1. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-APR-2000; 2000US-0200794P-
28-JUL-2000; 2000US-0221419P-
10-NOV-2000; 2000US-0247443P-
                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 14; Page 88; 239pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-075162/10.
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Ohmen JD, Reed
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 544 GTCGACTTTGAGGTGACA 561
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MONELL CHEM SENSES CENT.
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Tordoff MG;
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                                                                                                                                                              The invention relates to an isolated polypeptide, comprising a variant CC form of mouse or human SAC1 polypeptide. The variant form is associated CC with altered preference for carbohydrates, other sweeteners or ethanol. CC The polypeptide and its associated DNA sequence can be produced by CC recombinant techniques and is useful for preventing obesity, diabetes or clobolism associated with SAC1 expression. The sequences are useful in CC creening for drugs and sweeteners. Recombinant cell lines and transgenic cembryos may be used in screening for and identifying agents that induce CC or repress function of SAC1. Predisposition to diabetes, obesity or CC clobolism can be ascertained by testing any fluid or tissue of a human CC (such as blood, pancreas or tongue) for sequence variations of the SAC1 locus may indicate a CC diagnostic mark. The polynucleotide can be detected in a biological CC sample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and CC which is then detected. The sequences represent cDNA encoding human and CC mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes
                                                                   Matches
                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bachmanov AA, Beauc Ohmen JD, Reed DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; mouse; SAC1; carbohydrate; sweetener; ethanol; alcoholism; ss; obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas; blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy; protein replacement therapy.
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28-JUL-2000; 2000US-0221419P.
10-NOV-2000; 2000US-0247443P.
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                                                                                                                                       Sequence
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(MONE-) MONELL CHEM SENSES CENT.
                       6069 TAAATCTGGTTCTTTTTC 6086
18
                                                                   16;
                                                                                       Similarity
                                                                                                                                       20 BP; 9
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                                                                   Conservative
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d DR, Ross D,
                                                                                                                                       A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                  88.9%;
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                                                                                     Score 14.8;
Pred. No. 26
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Tordoff MG;
                                                                   Mismatches
                                                                                       2e+03;
                                                                                                     DB 1;
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RESULT 3034 AAS97784 ID AAS9778

AAS97784 standard; DNA; 20

RESULT 3035 AAS97786

AAS97786 standard; DNA;

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AAS97786

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                                                                                                                                             The invention relates to an isolated polypeptide, comprising a variant CC form of mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by crecombinant techniques and is useful for preventing obesity, diabetes or calcoholism associated with SAC1 expression. The sequences are useful in screening for drugs and sweeteners. Recombinant cell lines and transgenic embryos may be used in screening for and identifying agents that induce or repress function of SAC1. Predisposition to diabetes, obesity or calcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 (gene. A sequence variation of the SAC1 locus may indicate a predisposition to diabetes, obesity and/or alcoholism and may provide a diagnostic mark. The polymucleotide can be detected in a biological cample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequences represent cDNA encoding human and complex sequences sacc1 polypeptides and PCR primers specific for the SCA1 genes
                                                           Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
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28-JUL-2000; 2000US-0221419P
10-NCV-2000; 2000US-0247443P
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                                                                                                                       Sequence
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                              544 GTCGACTTTGAGGTGACA
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                                                                            Similarity
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                                                                                                                       BP;
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                                                            Conservative
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ed DR, Ross D,
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CHEM SENSES
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                                                         Pred. No. 2e+
0; Mismatches
                                                                                          Score 14.8;
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Tordoff MG;
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RESULT 3036
ABK37182/c
ID ABK3718
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AC ABK3718
XC ABK3718
XC ABK3718
XC O8-MAY-
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DT 08-MAY-
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DE Human 1
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                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                         182/c
ABK37182 standard;
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blood; tongue; PCR
                                               08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine SAC1 gene-specific oligonucleotide PCR
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10-NOV-2000;
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                                                                                                                                                                                                                                                                     544 GTCGACTTTGAGGTGACA
                                                                                                                                                                                                                                                                                                                               16;
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MONELL CHEM SENSES CENT.
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diabetes; transgenic embryo;
ondue; PCR primer; anorectic;
                                                                                                                                                                                                                                                                                                                                                                                                                  20 BP; 5
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ilarity 88.9%;
Conservative
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2000US-0221419P.
2000US-0247443P.
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                                               (first entry)
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1 DR, Ross D,
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                                                                                                                                DNA;
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Pred. No. 26
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Tordoff MG;
                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sweetener; ethanol; alcoholism; ss;
ryo; body tissue; body fluid; pancre
tic; antidiabetic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                 2e+03;
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Human lysophospholipase I gene,

antisense oligonucleotide #134

WO200236820-A2

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RESULT 3037
ABN83847/c
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Best Local S
Matches 16
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                                                 Non-insulin dependent diagnosis; prognosis; insulin; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel antisense compound useful for treating inflammation, hyperlipidemia, and cardiovascular disorders such as athen myocardial ischemia, inhibits Lysophospholipase I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; mouse; antiinflammatory; antiarteriosclerotic; vasot antilipaemic; cardiant; lysophospholipase I; inflammation; hyperlipidaemia; cardiovascular disorder; atherosclerosis;
                                                                                                      Insulin
                                                                                                                                  10-SEP-2002
                                                                                                                                                             ABN83847;
                                                                                                                                                                                         ABN83847
                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
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                                                                                                       MspI polymorphism PCR primer INS56
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                                                                                                                                                                                                                                                                                                                           0.2%;
                                                    88.
                                                              diabetes; type II diabetes; obesity; l
linkage disequilibrium; polymorphism;
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Pred. No. 26
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cion; ischaemia;
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                                                                marker;
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ABS74284/C
ID ABS7428
XX ABS7428
XX ABS7428
XX O9-DEC-
XX Human C
XX Human;
KW Gene th
KW autoimm
KW brain t
KW Korsa-k
KW Korsa-k
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KW SSCP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides methods for determining the risk of. CC development of non-insulin dependent diabetes mellitus (NIDDM or type II CC diabetes) in a subject. It results from the discovery that homozygotes of the HphI locus of the insulin gene along with body fat measurement serve conditions with HphI(+/-) or (-/-) genotypes are significantly more continued in the phI locus of the insulin HphI locus. The marker in linkage disequilibrium with the insulin HphI locus. The marker is especially -4217 PstI, -2221 MspI, -2231 MspI, +1428 PokI, +1000 AluI and +32000 ApaI. In an example, new polymorphisms were screened to determine whether they altered a controls, and a subset of polymorphisms was amplified using the primer INS57 to amplify the -2221 MspI site. PCR products were digested with MspI and gel electrophoresed to determine game yield using the primer INS57 to amplify the -2221 MspI site. PCR products were conducts of digestion were 108 and 78 bp for +/+, and 186 bp for -/- genotypes, where (+) indicates the restriction invention provides methods for diagnosting a subtype of NIDDM, for cestimating the frequency of a haplotype for a set of genetic markers in a population suffering from juvenile obesity (claimed), and methods to facilitate therapy and maintenance of NIDDM patients
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Best Local (
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Human; ss; primer; calcium channel alpha2delta; splice isoform; CACNA2D2; gene therapy; Lambert-Eaton myasthenic syndrome; LEMS; PCR; autoimmune disease; epilepsy; migraine; episodic ataxia; cancer; stroke; brain trauma; Alzheimer's disease; multiinfarct dementia; convulsion; Korsa-koff's disease; amyotrophic lateral sclerosis; seizure; Huntington's disease; amyotrophic lateral sclerosis; seizure; hypoxia; ischaemia; myocardial infarction; congestive heart failure; muscular dystrophy; hypertension; chromosome 3p2l.3; lung cancer; breast cancer; preneoplastic lesion; hyperplasia; dysplasia; carcinoma; SSCP; single strand change polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Determining risk of developing non-insulin dependent diabetes mellitus, comprises determining the identity of polymorphic bases of a marker in linkage disequilibrium with the insulin HphI locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-519258/55
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                                                                                                                                                                                                                                  calcium
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                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                  channel alpha2delta SSCP PCR primer #8.
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Pred. No. 2e
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RESULT 3039 ABA90030/c ID ABA9003 XX

ABA90030 standard; DNA; 20 BP

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20

Matches Query Match Best Local

16;

Conservative

0,

Mismatches

0

Gaps

0

2e+03;

Similarity

0.2%;

Score 14.8; Pred.

DB 1; Length 20;

5287 CAGCCTCTACTCCCAGCA 5304

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CC CACNADED gene which encodes human calcium channel alphadelta-2 subunit CC protein) comprising a fully defined alpha2delta splice isoform 1, 2 or 3 cc include are: (1) a method of producing a calcium channel proteins. Also cc include are: (1) a method of producing a calcium channel protein which cinvolves introducing a recombinant expression vector comprising the cc calculated host cell under conditions such that the host cell expresses the calcium channel protein, into a cc cultured host cell under conditions such that the host cell expresses the calcium channel protein, into a cc cultured host cell under conditions such that the host cell expresses the calcium one expression vector comprising one or more mucleic acid sequences encoding the splice variants. CACNAPD2 nucleic acid is useful for proteins, comprising carrying out the method of (1), but with one or more than one expression vector comprising one or more nucleic acid sequences composing the splice variants. CACNAPD2 nucleic acid is useful for polypeptide is useful for treating patients with Lambert-Eaton myasthenic compounds useful for treating other disease associated with abnormal calcium channel protein activity (e.g. epilepsy, migraine, episodic attaxia, cancer, stroke, brain trauma, Alzheimer's disease, multilinfarct dementia, carca-koff's disease, amyotrophic lateral sclerosis, convulsions, convulsions, the cardiovascular system, myocardial infarction, congestive heart calcium; muscular dystrophy and hypertension) CACNAPD2 nucleic acid is sequence encoding at least a portion of calcium channel protein, in detection, identification and isolation of alpha2delta sequences common in cancer (e.g. lung cancer and breast cancer) and served common in cancer (e.g. lung cancer and breast carcer) and served common in cancer (e.g. hupg cancer and breast carcer) and served common in cancer (e.g. hupg cancer and breast cancer) and content for the present is an SCCP (single strand change soluting a bruan calcium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 7; Col 46; 77pp;
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Sequence
                                                                             used to detect polymorphisms in sequences encoding channel alpha2delta splice isoform protein
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4 C; 10 G; 5 T; 0 U; 0 Other;
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ID AAD36736
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20-OCT-2000;
24-JAN-2001;
                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel variant of estrogen receptor alpha polypeptide useful for determining the biological activity of a protein for high throughput screening and for raising antibodies that elicit an immune response
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                 04-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-FEB-2001; 2001WO-US005358.
                                                       AAD36736;
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16; Conserv
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2000US-00692414.
2001US-00768184.
               (first entry)
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ABA97644/c
                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of diagnosing atherosclerosis or predicting the susceptibility of a subject to atherosclerosis, comprising determining the presence or absence of a single nucleotide polymorphism (SNP) in codon 379 of a lipoprotein-associated phospholipase A2 (Lp-PLA2) encoding polymucleotide, where the codon comprising the SNP encodes an amino acid other than valine. The method is useful for predicting the likelihood of developing atherosclerosis, predicting and responding to the progression of the atherosclerotic condition and drug treatment, or predicting disease outcome in a subject. The method is also useful in the selection of patient groups for conducting clinical trials concerning therapeutic compounds with potential for use in the treatment of atherosclerosis. The present sequence is human lipoprotein-associated phospholipase A2 gene amplifying primer used to detect polymorphism in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
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             Unidentified
                                      ss; fluorochrome;
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                                                                                                                 ABA97644;
                                                                                                                                       ABA97644 standard;
                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                    phospholipase A2 codon 379 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 9; Page 9; 30pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing atherosclerosis or predicting susceptibility of a subject to atherosclerosis by determining single nucleotide polymorphism in Lp-PLA
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/mod_base= OTHER
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                                      fluorescence
                                                                                                                                                                                                                                                                                                          U; 0 Other;
                                                                                                                                                                                                                                                                   2e+03;
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                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                         Gaps
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ABA97650/c
ID ABA9765
XX ABA9765
XX ABA9765
XX I1-APR-
XX INIDER Probe u
XX Unident
XX Unident
XX Unident
XX Unident
XX IF 20-APR-
PR 20-APR-
PR 20-APR-
PR 30-AUG-
PR (KANK-)
PA (KEIZ-)
XX WPI; 20
XX WPI; 20
XX Weasure
PT Measure
PT Measure
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Best Local
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30-AUG-1999;
01-FEB-2000;
                                                                                                                      20-APR-1999;
24-AUG-1999;
30-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a method for measuring nucleic acids using nucleic acid probe labelled with a fluorochrome. The nucleic acid probe control with a fluorochrome when hybridised with target nucleic acid, the decrease in the fluorescence is measured. I method can be used for measuring a target nucleic acid
                                                                                                                                                                                                                                                                                                                                              ABA97650 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 14 A; 1 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Measurement of nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-134193/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BIOI-)
(KANK-)
(KEIZ-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2001286300-A.
                                                                 (BIOI-)
(KANK-)
(KEIZ-)
                                                                                                                                                                                                                                                                                                    11-APR-2002
                                                                                                                                                                                                                                                                                                                          ABA97650,
                       Measurement of nucleic
                                                                                                                                                                                         16-OCT-2001
                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                         se; fluorochrome;
                                                                                                                                                                    20-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   obtained data.
             obtained
                                                                                                                                                                                                                                                                                                                                                                                                                         6682
                                                                                                                                                                                                                                                                                                                                                                                                   18
                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BIOINDUSTRY KYOKAI SH.
KANKYO ENG KK.
KEIZAI SANGYOSHO SANGY
                                                                BIOINDUSTRY KYOKAI SH.
KANKYO ENG KK.
KEIZAI SANGYOSHO SANGYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Page 18; 34pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                  TTATTTTTATTTATATAT 6699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; 99JP-00111601.
; 99JP-00236666.
; 99JP-00242693.
; 2000JP-00028896.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                            99JP-00236666.
99JP-00242693.
2000JP-00028896.
             data
                                                                                                                                                                    2000JP-00120097
                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                              99JP-00111601.
                                                                                                                                                                                                                                                        nucleic acid probe; fluorescence
                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acids, using
                       acids, using
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                                                                  GIJUTSU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GIJUTSU
                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           No.
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                       a nucleic acid probe and analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                           2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOGO
                                                                   SOGO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probe and analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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AAS16663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The
                                                                                                                                                                                                                                                                                                                                                                                                                              Human; inhibitor of DNA binding-1; Id-1; cycostatic; antiinflammationuppressive; antisense therapy; antisense oligonucleotide; hyperproliferative disorder; immune disorder; muscular disorder; vascular disorder; pancreatic disorder; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             method can
                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 18; 34pp; Japanese
                                                              Novel
                                                                                                                                                                                            08-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Inhibitor of DNA binding-1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS16663;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS16663
                    Example
                                         hyperproliferative,
                                                  expression
                                                                                   WPI; 2002-041477/05
                                                                                                        Baker
                                                                                                                                                 28-APR-2000;
                                                                                                                                                                      25-APR-2001;
                                                                                                                                                                                                                WO200183513-A2
                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                               modified
                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6682 TTATTTTTATTTATATAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
                                      . antisense compound, specifically hybridizing to and inhibiting the ssion of Inhibitor of DNA binding-1, useful for treating proliferative, immune, muscular, vascular or pancreatic disorder.
                                                                                                        BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                             SISI
                    15; Page 82;
                                                                                                                                                                                                                                                                                                             base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTTTTTTATATATATAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be used for measuring a target nucleic acid
                                                                                                        Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                       2001WO-US013209
                                                                                                                                                   2000US-00561497.
                                                                                                                             PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "2'-O-methoxyethyl
                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "Phosphorothioate backbone.
residues are 5-methyl cytidines"
                                                                                                                                                                                                                                     note=
                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                   16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 0 C; 0
                                                                                                                             INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
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                                                                                                                                                                                                                                     base= OTHER
e= "2'-O-methoxyethyl
                  105pp;
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                                                                                                       Wyatt
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                     English
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 5 T; 0 U; 0 Other;
                                                                                                        JR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide ISIS
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No. 2e+03;
                                                                                                                                                                                                                                      (2'-MOE)
                                                                                                                                                                                                                                                                               (2'-MOE)
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                                                                                                                                                                                                                                                                               nucleotides"
                                                                                                                                                                                                                                       nucleotides'
                                                                                                                                                                                                                                                                                                                                    Also,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiinflammatory;
                                                                                                                                                                                                                                                                                                                                    a11
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                                                                                                                                                                                                                                                                                                                                   cytidine
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The invention relates to novel antisense

compounds

(I) 8-30 nucleobases

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RESULT 3044
ABL94386/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                    LAP; TCF5; CRP2; NFIL6; IL6DBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness; oxidative stress response; IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                                              modified_base
                                                                                                                                                                    MUB
MUB
                                                                                                                                                                                                                                                                   Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta;
                                                                                              modified_base
                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                       Mouse
                                                                                                                                                                                                                                                                                                                               ABL94386;
                                                                                                                                                                                                                                                                                                                                                  ABL94386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                    musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                     3270
                                                                                                                                                                                                                                                                                      C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:152
                                                                                                                                                                                                                                                                                                                                                                                                                                        16;
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                                                                                                                                                                                                                                                                                                                                                                                                              ATTTGTTTAAGAAGAAAA 3287
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                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                   ATTTGTTTAATAACAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                          (first entry)
                                              cytosines
       /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE)
/rote= "2'-methoxyethyl (2'-MOE)
                                                       /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2'
cytosines are 5-methylcytosine"
                                                                                    /*tag= b
                                                                                                                                               Location/Qualifiers
                                                                                                       note= "Phosphorothioate linkages"
                                                                                                                   mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 C; 1 G; 9 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 14.8; DB No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1;
                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>ب</u>
                  nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                  A11
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                    C/EPB2;
                  ń
                  S
S
                                                                   SOM
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CC in length targeted to a nucleic acid molecule encoding Inhibitor of DNA C binding-1, where (I) specifically hybridises with and inhibits the CC expression of Inhibitor of DNA binding-1. Antisense inhibition of human CC Inhibitor of DNA binding-1 expression by chimeric phosphorothioate CC expression of DNA binding-1 expression by chimeric phosphorothioate CC oligonucleotides having 2' methoxyethyl (2'-MOE) wings and a deoxy gap CC was tested. A series of oligonucleotides were designed to target CC different regions of the human Inhibitor of DNA binding-1 RNA. The compounds were analysed for their effect on human Inhibitor of DNA binding-1 RNA. The CC monomial of the human Inhibitor of DNA binding-1 RNA. The compounds were sult showed that the oligonucleotides showed at least 25% CC inhibition of human Inhibitor of DNA binding-1 in cells or tissues by contacting the expression of Inhibitor of DNA binding-1 in cells or tissues by contacting the cells or tissues with (I). (I) is also useful CC for treating a human having a disease or condition associated with CC Inhibitor of DNA binding-1 by administering a therapeutically or CC prophylactically effective amount of (I), where the disease or condition is a hyperproliferative disorder, immune disorder, muscular disorder, CC diagnostics, therapeutics, prophylaxis (e.g., to prevent or delay condition sequence represents a human Inhibitor of DNA binding-1, antisense CC oligonucleotide used in the method of the invention binding-1, antisense CC oligonucleotide used in the method of the invention.
WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compound targeted to nucleic acids mouse CCAAT/enhancer binding protein (C/EBP) beta, inhibiting expression of human or mouse C/EBP beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 17; Col 49-50; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inhibiting expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUN-2000; 2000US-00593711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-00593711
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding human or useful in vitro for in cells/tissues.
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for

oligonucleotide used in the method of the invention

CC gene, which inhibit its expression. The antisense oligonucleotides were consigned to target different regions of the human and/or mouse C/EBP CC alpha RNA, and were analysed for their effect on C/EBP alpha mRNA levels by quantitative real-time PCR. The C/EBP family of proteins are a family CC of transcription factors which regulate the expression of a wide range of CC genes that control normal tissue development, cellular function, cellular CC proliferation and functional differentiation. C/EBP beta (also known as CC C/EBP2, LAP, TCP5, CRP2, NFIL6, IL6DBP, NF-M, ACP/EBP beta (also known as CC C/EBP2, LAP, TCP5, CRP2, NFIL6, IL6DBP, NF-M, ACP/EBP beta is continued to be involved in carbohydrate metabolism, immunity, the Thick thought to be involved in carbohydrate metabolism, immunity, the Thick expressed in the liver, lung, spleen, kidney, brain, and testis, with the highest expression found in the lung. It is also expressed at a higher CC level in malignant ovarian tissue compared with normal ovarian tissue, CC and its expression in pancreas is upregulated in response to chronically celevated levels of glucose, indicating that it is involved in the confidence of the invention are useful for diagnosis, prevention and treatment of conditions associated with C/EBP beta expression, such as cancer (particularly ovarian cancer), tumour formation, diabetes (particularly ovarian), rumour formation Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted the human or mouse CCAAT/enhancer-binding protein alpha (C/EBP alpha) H diabetes), infection, or inflammation

Sequence 20 BP; 0 Ŗ 9 C; 7 ဂ္ 4 T; ou; 0 Other;

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닭
           S
                            Query Match
Best Local S
Matches 16
              7414 AGCAGCAGCAGCAGC
18
                             16;
                                     Similarity
AGCGGCAGCAGCGCAGC 1
                             Conservative
                                   0.2%;
              7431
                             0
                                    Score 14.8;
Pred. No. 2
                             Mismatches
                                   2e+03;
                                           DB 1;
                             2;
                                            Length
                             Indels
                                            20;
                             0
                             Gaps
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0

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ABI93710
ID ABI9
RESULT 3045
                                           ABI93710 standard; DNA;
                                      ABI93710
                                            20
                                            BP
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Capture oligonucleptide Zip ID#797 oligo

16-FEB-2002

(first entry)

Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; Barl; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed

Synthetic

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RESULT 3046
ABI94356
ID ABI9435
XX
AC ABI9435
XY
I6-FEB-
XX
Capture
XX
Human;
KW lifacti
KW oncoger
XW enviror
XX
OS Synthet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cc for detecting infectious diseases caused by bacterial infectious agents ceg. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and ceg. Aspergillus fumigatus, viruses e.g. T-cell lymphocytotrophis cirus, cegetein-Barr virus and polio virus, and parasitic infectious agents cegetein-Barr virus and polio virus, and parasitic infectious agents cegetein-Barr virus and polio virus, and parasitic infectious agents cegetein-Barr virus and polio virus, and parasitic infectious agents cegetein-Barr virus and polio virus, and parasitic infectious agents cegetein-Barr virus and polio virus, and parasitic infectious agents cegetein-Barr virus endous security genetic diseases such cegetein-Barr virus endous security genetic diseases such cegetein-Barr virus endous security defects. The method is also useful for detecting genes, or genes certifically associated with a gene selected from BRCA1 gene, ps3 gene, human papillomavirus types 16 and 18 and liver cancers. The certification detecting comprises scanning forensics and the food and feed industry, detecting comprises scanning forensics and the food celectron microscope and infrared microscope) the support at the celectron microscope and infrared microscope) the support at the celectron cocurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. AB182074 to cef. The composition of the oligonucleotide sequences and the exemplification cef. for the careficiants of the composition to a presence of the target nucleotide sequences. AB182074 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                 ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious di
infection; 21 hydroxylase deficiency; Turner Syndrome; obesity;
oncogene; tumour suppressor; human papillomavirus; forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-APR-2001; 2001WO-US010958.
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                                               environmental
                                                                                                                                                                    Capture oligonucleptide Zip ID#1443 oligo #9
                                                                                                                                                                                                                  16-FEB-2002
                                                                                                                                                                                                                                                                                                ABI94356 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    have melting temperatures within a narrow range. The method is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16;
                                                                                                                               K-ras;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                             monitoring; food
                                                                                                                       PCR primer; probe; capture probe; mutation detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8;
Pred. No. 26
                                             industry; feed industry; ss
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CC oligonuclectide probes (II) will hybridise with little mismatch, where (II) have melting temperatures within a narrow range. The method is useful (II) have melting temperatures within a narrow range. The method is useful (II) detections diseases caused by bacterial infectious agents (II) will hybridise and harmow range. The method is useful (II) fectious agents e.g. Cryptococcus meoformans, Candida albicans and (II) septeillus funiquitus, viruses e.g. T-cell lymphocytotrophils cirus, (II) selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus (II) medinesis. The method is also useful for detecting genetic diseases such (II) detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the (II) gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (Using e.g. a scanning (II) september of the support at the particular sites and infrared microscope) the support at the probe sets occurred and correlating (Using a computer) identified ligation to a gressance or absence of the target nucleotide sequences. ABI92074 to of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention describes a method (M1) for designing capture ride probes (I) for use on a support to which complementary
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Sequence 20 BP; 3 A; c; G; 6 T; 0 U; 0 Other;

S Matches Query Match Best Local (3117 TGCTTGACAGCTTGGTAA 3134 16; Similarity Conservative 0.2%; ç, Score Pred. Mismatches e 14.8; b. DB 1, Length Indels 20; 0; Gaps 0

ABK28754 standard; DNA; 20 ВP

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Human CDC14 gene donor splice site #14.

09-APR-2002 ABK28754;

(first entry)

Human; ds; cell-cycle control; CDC14A; cancer; splice site; prostate cancer; breast cancer; tumour; lymph node metastasis; malignant mesothelioma; chromosome 1p21; dual specificity phosphatase; therapy; replacement therapy

Homo sapiens

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ABL56505/A

ABL556505/A

AC ABL55

AC ABL55

AX ABL50

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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an isolated nucleic acid encoding a CDC14A polypeptide (cell-cycle control protein 14A, a dual specificity phosphatase), its complement or RNA molecule corresponding to it. Also included are an expression vector comprising the nucleic acid and a host cell transformed with the vector. The gene for CDC14A is located on human chromosome 1p21. The nucleic acid and protein are useful to diagnose and treat human cancers (e.g. breast cancer, prostate cancer) and tumours (e.g. lymph node metastasis, malignant mesothelioma) which have a mutation in the CDC14A gene, by gene therapy, protein replacement therapy or protein mimetics. They can also be used to screen for drugs to treat sequence is a splice donor or splice acceptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid encoding mutated form of human dual-specificity phosphatase CDC14A polypeptide, useful to diagnose and treat cancers.
    WPI; 2002-340014/37
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                                                   Waksman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABL56505 standard;
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16; Conserv
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                                                   Lamartine
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Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                                              Clouston syndrome; alopecia; hair loss; hair growth; hair disorder; PCR; primer
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2e+03;
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ABQ87739/c
ID ABQ87739 standard; DNA;
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24-JAN-2001;
13-MAR-2001;
05-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primers ABL56504-07 were used to amplify fragments of the human GJB6 gene. The GJB6 gene encodes connexin-30 (Cx-30). A G to A mutation at position 31 and/or a C to T mutation at position 263 causes Clouston syndrome (hydrotic ectodermal dysplasia). Nucleic acids encoding Cx-30, antisense oligonucleotides and vectors containing them, are used to treat Clouston syndrome and other forms of alopecia with a genetic component. They are also used to reduce growth of hair and/or promote hair loss. They may also be used for (cosmetic) treatment and/or prevention of disorders of the hair. Transgenic animals/cells that contain Cx-30 are used to screen for agents that are potentially useful for treating some forms of alopecia. Primers derived from GJB6 gene are useful in
                                                                                                                                               Kalush
                                                                                                                                                                                                                                                                         02-MAY-2002.
                                                                                                                                                                                                                                                                                                WO200234945-A2
                                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                            ESR-alpha; ESR1;
                                                                                                                                                                                                                                                                                                                                                       Human; oestrogen; rece; osteopathic; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New mutant for
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                                                                                                                                                                                                                                                                                                                                                                                        Human ESR1 exon 8.18 sequencing PCR primer ER1x8.18sr3_54048.
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                                                                                                                                                                     (APPL-) APPLERA CORP
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; 2001US-00804076.
; 2001US-00826314.
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Clouston syndrome
                                                                                                                                                                                                                                                                                                                                            sequencing; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                receptor; oestrogen receptor
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                                                                                                                                                                                                                                                                                                                                                        cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8;
Pred. No. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 T;
                                                                                                                                                                                                                                                                                                                                                       osteoporosis; cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid, useful for the diagnosis ome and other forms of alopecia.
                                                                                                                                           Winn_deen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 U;
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                                                                                                                                                                                                                                                                                                                                                        disorder;
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The invention relates to novel human oestrogen receptor variant peptides, and the polynucleotides encoding them. The peptides of the invention have cytostatic, osteopathic and cardiant activity. The peptides of the

mediate or modulate

Peptide of estrogen receptor alpha genes variant or its in identifying modulators for treating disorders e.g. a

cardiovascular

disorder.

fragment for use susceptibility to

Example

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Fig

2E; 352pp;

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osteoporosis,

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AAL47461
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Best Local Similarity
Matches 16; Conserv
                                                   Query Match
Best Local
                                         Matches
                                                                                                                        The present invention relates to a method of determining a nucleic acid sequence, involving incubating with a probe, adding at least one phosphorothioate nucleotide in presence of an enzyme that synthesises a complementary sequence on the 3'-end of the probe, and adding a second enzyme that degrades the phosphorothioate-free complement. The method can be used to determine very short (up to 10 base pair) sequences, especially for characterisation of single-nucleotide polymorphisms. The present sequence is a probe for the human MTHFR gene which was used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as a susceptibility to cancer, osteoporosis, cardiovascular disorder, etc., and hence are useful in the treatment of the disorders. The sequences shown in ABQ87720-ABQ87746 represent PCR primers used in the invention to sequence individual exons of the human oestrogen receptor
                                                                                       Sequence
                                                                                                                                                                                                                                                                     Determining nucleic acid sequence, useful for characterizing single-
nucleotide polymorphisms, by incubating with probe in presence of
phosphorothioate nucleotide and exonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                 20-NOV-2000; 2000DE-01058915
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human MTHFR gene probe.
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                                                                                                                  demonstrate the
                                                                                                                                                                                                                                                                                                                                                (ATTO-) ATTOMOL GMBH
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5816 CTATGTGATGATGAAATC
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                                       l Similarity
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ilarity 88.9%;
Conservative
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                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequencing; single nucleotide polymorphism; SNP;
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                                                                                                                  method
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'note= "modified by FITC"
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                                                 0.2%;
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                                                                                                                  of the invention
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pred. No. 2e+0
0; Mismatches
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Pred. No. 2e+03;
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RESULT 3051
RAL49188/c
ID AAL4918
XX AAL4918
XX AAL4918
XX AAL4918
XX AAL4918
XX AAL4918
XX CD 151;
XW CD 151;
XW Eslecti
XX CD 151;
XW Selecti
XX CD 151;
XW Selecti
XX CD 151;
XW SOLO 861
XX CD 151;
XX CD 151
RESULT 3052
ABZ92414
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                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method of determining the susceptibility or resistance of an animal to porcine reproductive and CC respiratory syndrome virus (PRRSV). This involves assaying for CD 151 in C a sample of cellular material of known origin from the animal. In CC addition, coding sequences of CD 151 are described, and anti-viral CC compounds designated anti-RNA entry proteins (anti-REPs). The method is C useful for determining susceptibility and resistance to PRRSV in an CC animal. This is particularly useful for improving swine breeding or for CC screening different pig breeding lines. The method is also useful for CC certains on-simian recombinant cell lines for propagating the virus, CC developing non-simian recombinant cell lines for inducing immunity CC against PRRSV, and for diagnosing PRRSV infection in a swine. The present CC sequence is a PCR primer used to isolate the porcine CD 151 coding CC the printed specification, but was obtained in electronic format directly CC from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Determining susceptibility and resistance to porcine reproductive and respiratory syndrome virus (PRRSV), useful for improving swine breeding, by assaying for CD 151 in a sample of cellular material of known origin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD 151; porcine reproductive and respiratory syndrome virus; PRRSV; pig; selective breeding; xenotransplant; anti-RNA entry protein; anti-REP; anti-viral; vaccine; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JAN-2001; 2001US-00772044.
28-JAN-2002; 2002US-00772044.
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                                                                                                                                                                                                                                                                                                                                                              Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-2002; 2002WO-US002868.
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                                                                                                                                                                 993 CAAGGGCCTGAAGGTGGA 1010
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                                                                                                                                                                                                                                                                                                                                                              BP; 2 A; 7 C; 4 G; 7 T; 0 U;
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ABZ92414

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                                  RESULT 3053
ABZ92635/c
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CC Note: The sequence data for this patent is not represented in the printed cat firm who, intronchance of contraction, but was obtained in electronic format directly from WIPO cat firm, who, intronchance of the composition of the printed cat for this patent is not represented in the printed cat firm who intronchance is contented.
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, antisense; lung dysfunction; nasal airway dysfunction; antialflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                  ABZ92635
                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-APR-2001; 2001US-0286137P
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                                                                                                                                                                                                                                                                                                    ftp.wipo.int/pub/published_pct_sequences
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₹S,
                                                                                                                                                   3226
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                standard; DNA;
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Tang
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Conservative
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և, Shahabuddin S;
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RESULT 3054 ABZ88435 ID ABZ8843

ABZ88435 standard; DNA; 20

ВP

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TTCTAAGCAGATTGGGAA 6812

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Best Loc Matches

Local

Similarity

88.9%;

Pred.

No.

2e+03;

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Gaps

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 3' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of antiinflammatory steroid in a subject, for reducing levels of adenosine CC receptor, producing bronchodilation, increasing levels of adenosine CC lung surfactant in a subject stissue, or treating bronchoconstriction, CC lung inflammation, lung allergies, or a respiratory disease or condition. CC specification, but was obtained in electronic format directly from WIPO or after wino int/mih/shad not semences.
 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                       Sequence
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antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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Tang L,
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                                     BP; 6 A;
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L, Shahabuddin
                                   5 C;
0.2%;
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 Score 14.8;
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DB 1; Length 20;
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RESULT 3055
ABZ90375/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
               ABZ90375
                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 7 C; 1 G; 9 T; 0 U; 0 Other;
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                                                                                                                                                  2167 TACAAGTCCACCCGTTTC
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                                                                                                                                                                                                       Similarity
               standard; DNA;
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Tang L,
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L, Shahabuddin
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ABZ87286 ID ABZI XX

ABZ87286 standard; DNA; 20

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RESULT 3056

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The invention relates to a novel pharmaceutical composition, which has a cfirst active agent comprising an oligonucleotide antisense to the cinitiation codon, coding region, 5' or 3' end genomic flanking regions, cc 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of cc junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an cc antiinflammatory steroid and ubiquinone. A composition of the invention cc antiinflammatory, antiallergic, antisthmatic, hypotensive, cc immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cf enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of biquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, cl ung inflammation, lung allergies, or a respiratory disease or condition. Cc lung inflammation, but was obtained in electronic format directly from WIPO at the payment directly from WIPO at the printed specification, but was obtained in electronic format directly from WIPO cr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                                     Sequence 20 BP; 16 A; 4 C; 0 G; 0 T; 0 U; 0 Other;
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                                    4464 TTTTTTTTTTTTTTTT 4481
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L, Shahabuddin
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RESULT 3057
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Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                ABZ87732
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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              standard; DNA;
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Tang
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Conservative
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L, Shahabuddin
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ABZ93605/c ID ABZ936

ABZ93605 standard; DNA; 20

ВP

RESULT 3058

문 8

19

Best Loc Matches

Local

Similarity

88.9%;

Pred. No.

2e+03;

<u>,,</u>

Mismatches

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Gaps

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2999

CCCCACCCCTCACCCCAT 3016

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or CC preventing a respiratory, lung or malignant disease or condition, also CC of or reducing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels CC antiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine cof lung surfactant in a subject's tissue, or treating bronchoconstriction, CC lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed CC specification, but was obtained in electronic format directly from WIPO CC at fig. vipo.int/pub/published_pct_sequences
Query Match
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Miller S,
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antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;
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antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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, Tang L,
                                       BP; 1 A; 1 C; 15
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L, Shahabuddin
0.2%;
                                       G; 3
  Score 14.8;
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                                       T; 0 U; 0 Other;
DB 1; Length 20;
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                                                                                                         RESULT 3059
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                                                                      ABZ97793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cy junctions of genes encoding a polypeptide associated with lung and/or compasal airway dysfunction and a second active agent comprising an compastinifiammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, has antiinflammatory, antiallergic, antiasthmatic, hypotensive, communosuppressive, and cytostatic activity. The composition may have a composition may have a composition antisense gene therapy. The composition is useful for treating or compressive, antistratory, lung or malignant disease or condition, also composition as a subject, for reducing or depleting levels of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine composition, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of adenosine composition in a subject stissue, or treating bronchoconstriction, composition inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed general composition, but was obtained in electronic format directly from WIPO as the print of the 
                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or corresponding transfer in the corresp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel pharmaceutical composition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 8847; 872pp; English.
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                                   ABZ97793 standard;
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, Tang L,
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L, Shahabuddin
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                                   DNA;
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Pred. No. 2e
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ABZ93220/ ID ABZ9 XX

ABZ93220 standard; DNA;

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RESULT 3060

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Matches
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                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Miller
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                 2405 GGACCACAGTGGACACCA 2422
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                                                        Conservative
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L, Shahabuddin S;
                                                                      88.9%;
                                                                                      0.2%;
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ABZ86063/c
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Matches 16
                     ABZ86063 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                         Sequence
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Tang
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                  T; 0 U; 0 Other;
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ABZ90434 standard; DNA; 20

RESULT 3062 ABZ90434

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7415 GCAGCAGCAGCAGCA 7432

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Local

Similarity

0.2%;

Score 14.8; Pred.

DB 1; Length

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Mismatches

Indels

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Gaps

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of gunctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmattic, hypotensive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC constanting a respiratory, lung or malignant disease or condition, also CC for reducing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels CC antiinflammatory steroid in a subject, for reducing evels of adenosine coffice of the sequence of this patent is not represented in the printed CC Note: The sequence data for this patent is not represented in the printed CC specification, but was obtained in electronic format directly from WIPO CC at fig. wipo.int/pub/published_pct_sequences
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    Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nyce JW,
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, Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID
BP; 2 A; 5 C; 7 G; 6 T; 0 U;
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L, Shahabuddin
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    0 Other;
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RESULT 3063
ABZ90042
ID ABZ9004
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Note: The sequence data for this patent is not represented in the printed appectification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                    Query Match
Best Local S
Matches 16
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                    ABZ90042 standard; DNA;
                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human oligonucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel pharmaceutical composition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 5676; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-229219/22.
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                                                                                                                                                      1341 GATCAGTCGCCTGATGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation; respiratory disease; ds.
                                                                                                                         L
                                                                                                                                                                                                                        Similarity
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, Tang L,
                                                                                                                                                                                                                                                                                  20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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L, Shahabuddin
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                      20 BP
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Pred. No. 26
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ABZ85315

standard; DNA; 20

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RESULT 3064 ABZ85315/c

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The invention relates to a novel pharmaceutical composition, which has a composition codon, coding region, 5 or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 6' or regions within 2-10 nucleotides of 6' junctions of genes encoding a polypeptide associated with lung and/or 6' or comparison of the invention 6' or assal airway dysfunction and a second active agent comprising an 6' or assal airway dysfunction and a second active agent comprising an 7' or composition at a subject, antiasthmatic, hypotensive, 7' or immunosuppressive, and cytostatic activity. The composition may have a 7' or enthancing the prophylactic or the composition is useful for treating or 7' or preventing a respiratory, 1' ung or malignant disease or condition, also 6' or enhancing the prophylactic or therapeutic respiratory effect of an 7' or reducing the prophylactic or therapeutic respiratory effect of an 7' or reducing sensitivity to adenosine, reducing levels of adenosine 7' or reducing bronchodilation, increasing levels of ubiquinone or 8' or 1' or 1'
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                                                                                            Matches
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Miller S
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                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 5284; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-APR-2001; 2001US-0286137P
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                                                                                                                Local
                          3955 TCTTATGTTTCAATATTT 3972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide sequence.
                                                                                            16;
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                                                                                                                Similarity
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, Tang L,
TCTTATGTTTCTAAATTT
                                                                                                                                                                                    BP; 4 A; 2 C; 2 G; 12 T; 0 U; 0 Other;
                                                                                            Conservative
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L, Shahabuddin
                                                                                                                88.9%;
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                                                                                          0,
                                                                                                                     Pred.
                                                                                                                                          Score 14.8;
                                                                                            Mismatches
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                                                                                                                     No.
                                                                                                                2e+03;
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ABZ93982/c
ID ABZ939
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                                                                                                                                                                                                                                                        CC The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5 or 3 end genomic flanking regions, CC 5 and 3 intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels of of or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or CC lung surfactant in a subject's tissue, or treating bronchoconstriction, CC Note: The sequence data for this patent is not represented in the printed Specification, but was obtained in electronic format directly from WIPO CC at fig. wipo.int/pub/published_pct_sequences
                                                   RESULT 3065
                                                                                                                                                                      Matches
                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
               ABZ93982 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antia
                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
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                                                                                                                                     6461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15; SEQ ID NO 557; 872pp; English.
                                                                                                   19
                                                                                                                                                                                     Similarity
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                                                                                                                                 ATACTTTTTTTTTCTGTT 6478
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                                                                                                                                                                                                                                      B₽;
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                                                                                                                                                                                                                                      12
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L, Shahabuddin
                                                                                                                                                                                                                                     A; 3 C; 2
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                                                                                                                                                                                   0.2%;
                                                                                                                                                                     0
                                                                                                                                                                                                                                     G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                     Score 14.8;
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                                                                                                                                                                     Mismatches
                                                                                                                                                                                     2e+03;
                                                                                                                                                                                                   DB 1; Length 20;
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                                                                                                                                                                     Gaps
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RESULT 3066 ABZ98928/c ID ABZ98928 & XX

standard; DNA; 20

밁 S

20

Matches Best Local Query Match

Similarity

88.9%; 0.2%;

Score 14.8; Pred.

DB 1; Length 20;

Conservative

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Mismatches

0

Gaps

0

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3302 AGATCAATATTTTAGGAT 3319

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC compositing a respiratory, lung or malignant disease or condition, also CC containing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels of an containinflammatory steroid in a subject, for reducing adenosine coft or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine coft on surfactant in a subject's tissue, or treating bronchoconstriction. CC lung surfactant in a subject's tissue, or respiratory disease or condition. CC where the sequence data for this patent is not represented in the printed conscription, but was obtained in electronic format directly from WIPO CC at fig. vipo.int/pub/published_pct_sequences
Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-APR-2001; 2001US-0286137P
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, Tang L,
BP; 7 A; 6 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 9224; 872pp; English
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L, Shahabuddin
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RESULT 3067
ABZ88693/c
ID ABZ8869
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                                                                                                                                                                                                                                                                                                            The invention relates to a novel pharmaceutical composition, which has a confirmation for the segment comprising an oligonucleotide antisense to the confirmation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or antiinflammatory steroid and ubiquinone. A composition of the invention cathinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a composition is useful for treating or greventing a respiratory, lung or malignant disease or condition, also composition for the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine code subject, and subject at tissue, or treating bronchoconstriction, lung surfactant in a subject at tissue, or treating bronchoconstriction. CC lung inflammation, lung allergies, or a respiratory disease or condition. Subjective trissue, or treating bronchoconstriction, but was obtained in electronic format directly from WIPO care for which in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                       Best Local
Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cyrostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity;
                   ABZ88693 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 14170; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ubiquinone.
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adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PDE4A oligonucleotide sequence.
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                                                                                                                                                  6406
                                                                                                                 19
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                                                                                                                                          CCTGCTAGATAGCTTCTC
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                                                                                                                                                                                                                                                               BP; 6
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Conservative
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L, Shahabuddin
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0; Mismatches
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ABZ91491 standard; DNA; 20

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RESULT 3068 ABZ91491/c

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                                                                   Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW,
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                                                                                                                                       Sequence 20
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                                   5473
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18
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                                                                                      Similarity
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Tang
 TTTTTTTTAAAAAAATA
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L, Shahabuddin
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                                                                                                         Score 14.8;
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RESULT 3069
ABZ93213/c
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                       preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
               3213/c
ABZ93213 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 6733; 872pp; English.
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                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                        ВP;
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ilarity 88.9%;
Conservative
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L, Shahabuddin S;
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                                                                                                                                                                                                Pred.
                                                                                                                                                                                                                   Score 14.8;
                                                                                                                                                                                                                                                        G; 3
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RESULT 3070 ABZ92117 ID ABZ9211 XX

ABZ92117 standard; DNA;

20 ВP

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of continuing sensitivity to adenosine, reducing evels of adenosine CC for reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of adenosine of lung surfactant in a subject's tissue, or treating bronchocoonstriction. CC lung surfactant in a subject's tissue, or respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed conscription of the printed specification, but was obtained in electronic format directly from WIPO can first the printed specification in the printed specification in the printed specification.
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                                                                                Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nyce JW,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; lung dysfunction; nasal airway dysfunction;
antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ubiquinone.
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                   Госат
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                   Similarity
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Tang L,
                                                                                BP; 11 A; 2 C; 5
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L, Shahabuddin
                 0.2%;
0;
                   Pred.
                                         Score 14.8;
                                                                                G; 2 T; 0 U; 0 Other;
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    Mismatches
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                     No.
                 2e+03;
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                                 RESULT 3071
ACC62225/c
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                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel pharmaceutical composition, which has a confirmation codon, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding and a coding regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or coding and a second active agent comprising an continflammatory steroid and ubiquinone. A composition of the invention and a second active agent comprising an communication of the invention coding and cytostatic activity. The composition may have a communosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also confiniflammatory steroid in a subject, for reducing or depleting levels of antisinflammatory steroid in a subject, for reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or composition and antising antisort's risasse or readers of ubiquinone or composition and composition are composition.
                                                                                                                                                                                       Matches
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Best Local
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Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                 ACC62225
                                                                                                                                                                                                                                                                                                                  lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200285308-A2
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                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                4301 TCTTTTTCCTTCCCCTGG 4318
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                                                                                                             N
                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                               20
                 standard;
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                                                                                                                                                                                                                                                             BP; 0
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L, Shahabuddin
                                                                                                                                                                                                                                                             A; 9
                 DNA;
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                                                                                                                                                                                     Score 14.8; I
Pred. No. 2e+0
0; Mismatches
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The invention relates to a novel compound that is 8-50 nucleotides in CC length that is targeted to a nucleic acid molecule encoding apolipoprotein B (ApoB), and specifically hybridises with and inhibits CC specifically hybridises with at least an 8-nucleotide portion of an active site on a nucleic acid molecule encoding ApoB; or which CC specifically hybridises with at least an 8-nucleotide portion of an CC anterior has antilipaemic, antiarteriosclerotic, antidiabetic, and cardiovascular activity. The compound may have a use in CC gene therapy. The antisense oligonucleotide is useful for treating an CC condition involving abnormal lipid metabolism, a condition involving abnormal lipid metabolism, a condition involving CC abnormal cholesterol metabolic condition (e.g. hyperlipidaemia, diabetes (geocifically Type 2 diabetes), obesity, atherosclerosis or cardiovascular disease). The new compound or the antisense CC (specifically VLDL, HDL or LDL) or serum triglyceride levels (particularly in a human. The antisense compound or the ansociated for particularly in a human. The antisense compound or the associated with ApoB, or the onset of a disease or condition of the exemplification of the animal or the ansecoiated with ApoB, or the onset of an increase in glucose levels in the animal or the annimal or the animal o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              anorectic; cardiovascular; gene therapy; lipid metabolism; cholesterol metabolism; atherosclerosis; hyperlipidaemia; type 2 diabetes; obesiden, atherosclerosis; cardiovascular climose.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides for modulating apolipoprotein B, especially for preventing or treating atherosclerosis, hyperlipidiables, or for modulating glucose, cholesterol, lipoprotein or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-AUG-2001; 2001US-00920033.
30-APR-2002; 2002US-00135985.
15-MAY-2002; 2002US-00147196.
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anorectic; cardi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACC62225;
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Query Match
Best Local Similarity
Matches 16; Conserv

0.2%;

Score 14.8; D Pred. No. 2e+0 0; Mismatches

2e+03;

DB 1; Length <u>ب</u>

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6291 ACACTGGCCTCCAGGGAT

6308

Sequence 20

B₽;

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cc a bovine acyl CoA-diacyJglycerol transferase (DAT) contributing to or conditative for low fat content of milk and to low meat marbling contributing to or contributing the gene contribution that content of milk and/or its contribution for its predisposition for fat content of milk and/or its contributing. The method comprises analysing the gene contributing that content of milk and/or its contribution contributing. The method comprises analysing the gene contribution contribution contribution. The contribution contribution
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                                                                                                                                                                       AKEKABKKARKROKEKEKEKEKEKEKEFEEKKKOOOOOOOOOOOOOO
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ABZ77000
                                                                                 Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Acyl CoA:diacylglycerol transferase; DGAT; emilk; meat marbling; low fat; polymorphic; single nucleotide polymorphism; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 36; 91pp;
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13-MAY-2002; 2002US-0379412P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bovine DGAT PCR primer #36.
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16; Conserv
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                                                                                 Conservative
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18
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Pred. No. 2e+0
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                                                                                                     2e+03;
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SNP;
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                                                                                                                       Length 20;
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Matches Query Match Best Local :

16;

Conservative

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Mismatches

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Similarity

0.2%;

Score 14.8; Pred. No. 26

2e+03;

DB 1; Length

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Sequence 20

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Other,

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RESULT 3073
ABZ76933
CC a bovine acyl CcA-diacylglycerol transferase (DGAT) contributing to or contributing to or contributing countributing and covine DGAT is located to chromosome 8, and countributing and countributing countributing and countributing countrib
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a nucleic acid molecule (NA) (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid molecule comprising a sequence of an allele of a polymorphic bovine acyl CoA-diacylglycerol transferase gene useful testing a mammal for its predisposition for fat content of milk and meat marbling.
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13-MAY-2002;
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2002US-0379412P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymorphism; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       low fat; polymorphic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 #6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enzyme;
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ARESULT 3074
ART34193/c
ID ART34193/c
XX ART3415
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XX ART3415
XX MOUSE &
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KW Antiart
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RESULT 3075
ABZ70582/c
ID ABZ7058
XX
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Best Local :
                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel compound of 8 - 50 nucleobases in length targeted to a nucleic acid molecule encoding a short heterodimer partner-1. The novel compound specifically hybridizes with a nucleic acid molecule encoding the short heterodimer partner-1, and inhibits the expression of the nucleic acid molecule. The compound, and a composition comprising it are useful for treating a disease or condition associated with the short heterodimer partner-1, particularly a condition involving abnormal lipid or cholesterol metabolism such as atherosclerosis or a cardiovascular disease. They are also useful in research and diagnostics for modulating the expression of short heterodimer partner-1. They can also be useful prophylactically in preventing or delaying infection, inflammation or tumour formation. This polynucleotide sequence represents a mouse antisense oligo relating to the heterodimer partner-1 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antiarteriosclerotic; cardiant; vasotropic; antiinfective; cytostatic; antiinflammatory; inhibitor; antisense gene therapy; atherosclerosis; short heterodime; partner-1; abnormal; lipid; cholesterol metabolism; cardiovascular disease; infection; inflammation; tumour formation; mouse;
                            ABZ70582 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotide targeted to a nucleic acid encoding short heterodimer partner-1, useful for treating diseases involving abnormal lipid or cholesterol metabolism, e.g atherosclerosis or cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-248161/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 16;
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                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                               ATGTCTCAGCCACTGCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          heterodimer partner-1
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                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4
                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                           3 C; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                               G; 4 T; 0
                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1; Length 20; Pred. No. 2e+03; o; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression oligo
                                                                                                                                                                                                                                                                                                                                                                                                                           U; 0 Other;
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88X48X8X4XXX

antisense;

Homo sapiens

Apolipoprotein(a); antiarteriosclerotic;

cardiant; gene therapy; human;

Human apolipoprtein(a) mRNA inhibiting antisense oligo ISIS 144373.

ACC47290

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11-AUG-2003

(first entry)

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RESULT 3076
ACC47290/c
ID ACC47290 standard; DNA;
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                                                                                                                                                 Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Determining the risk of developing obesity for treating or reducing the risk of developing obesity by determining a paternal insulin variable number of tandem repeats allele in the individual.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-JUL-2001; 2001US-0309235P.
21-AUG-2001; 2001US-0316830P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 16; 62pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Insulin; genotyping; obesity; variable number of tandem repeats; VNTR;
human; PCR; primer; ss.
                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-248187/24.
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                                                                                                     4729 CTTGGAGGCCAGCTGGAG 4746
                                                                                    18
                                                                                                                                                   16;
                                                                                                                                                                   Similarity
                                                                                    CTTGAAGGCCAGCTGGTG 1
                                                                                                                                                                                                                 BP; 4 A; 7 C; 5
                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VNTR allele genotyping primer INS56.
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                                                                                                                                                                                                                   G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                    Score 14.8;
                                                                                                                                                                    Pred.
                                                                                                                                                     Mismatches
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RESULT 3077
ABX04329/c
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ABX0432
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AC ABX0432
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AC A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a new compound, 8-50 nucleobases in length targeted to a nucleic acid molecule encoding human apolipoprotein(a), specifically hybridizes with and inhibits the expression of human apolipoprotein(a). The antisense compounds are useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis or cardiovascular disease. Sequences ACC47284-318 represent specific examples of chimeric antisense phosphorothicate oligonucleotides having 2'-MOE wings and a deoxy gap targeting human onligonucleotides having 2'-MOE wings and a deoxy gap targeting human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
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                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
          Dean NM,
                                                                                                                                                                            26-MAR-1999;
17-MAR-2000;
                                                                                                                                                                                                                                                  07-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse; ss; antisense;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-2001;
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                                                                             DEAN N M.
KARRAS J G.
MCKAY R.
                                                       MANOHARAN M.
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       Karras JG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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2000WO-US007318
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       Mckay
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Pred. No. 2e+0;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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ABX17722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      versus soluble form of IL-5 receptor a. The antisense compound is useful for modulating IL-5 signal transduction, modulating expression of mammalian IL-5 or modulating the expression of mammalian IL-5 receptor a, in cells or tissues, for altering the ratio of the isoforms of mammalian IL-5 receptor a in mammalian cells or tissues, treating a mammalian having a disease or condition associated with IL-5 signal transduction, IL-5 expression or IL-5 receptor a expression, where the disease or condition include eosinophilic syndrome or asthma. An antisense compound which alters splicing of an RNA encoding IL-5 receptor a is also useful for treating a mammal having a disease or condition. The present sequence is an antisense oligonucleotide targetting mouse IL5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an antisense compound of 8-30 nucleobases in length, which modulates interleukin (II)-5 signal transduction. Also include are a pharmaceutical composition comprising the antisense oligonucleotide and a pharmaceutically acceptable carrier or diluent, and a diagnostic kit for detecting the expression level of the membrane form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI:, 2003-039602/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel antisense compound for treating disease/condition e.g. eosinophilic syndrome or asthma associated with interleukin-5 or IL-5 receptor expression or IL-5 signal transduction, modulates IL-5 signal
                                                                                                                                              30-MAR-2001;
                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                               antisense
                                                                                                                                                                                                                                                                                                                bone cancer; brain cancer; ovary can
endometrium cancer; stomach cancer;
                                                                                                                                                                                                                                                                                                                                                          Urokinase plasminogen activator; gene therapy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                      Human urokinase plasminogen activator antisense oligonucleotide #27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABX17722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABX17722 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 14; Page 16;
                                                                                    Baker BF,
                                                                                                                                                                           18-MAR-2002; 2002WO-US008112.
                                                                                                                                                                                                         10-OCT-2002.
                                                                                                                                                                                                                                       WO200279515-A1
                                                                                                                                                                                                                                                                                                                                           hyperproliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                       05-FEB-2003
                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5272 ATAGGGAGCAGGTGGCAG 5289
                                                        2003-058441/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 AGACGGAGCAGGTGGCAG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                    Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9 C; 3
                                                                                    Watt
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                                                                                                                                                                                                                                                                                                 88.
                                                                                                                                                                                                                                                                                                                             ovary cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G; 5 T;
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Pred. No. 26
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U;
                                                                                                                                                                                                                                                                                                                kidney cancer; tumour metastasis;
                                                                                                                                                                                                                                                                                                                                           breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2e+03;
                                                                                                                                                                                                                                                                                                                              cervix cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                              cancer;
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                                                                                                                                                                                                                                                                                                                                            colon cancer;
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New antisense compound, useful for preparing a composition for treating hyperproliferative disorders, cancer e.g., breast, colon, bone, brain, ovary, cervix, endometrium, stomach or kidney cancer, or tumor

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RESULT 3079
AAD53839/c
ID AAD5383
XX AAD5383
XX AAD5383
XX AUPRIA
XX Juvenil
KW Juvenil
KW Jivenil
AXX Jivenil

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Best Local (
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A new compound, which is 8-50 nucleobases in length targeted to a nucleoic acid molecule encoding urokinase plasminogen activator, specifically hybridises with and inhibits the expression of urokinase plasminogen activator. The compound is useful for preparing a composition for treating (e.g. by gene therapy) hyperproliferative disorder, cancer e.g., breast, colon, bone, brain, ovary, cervix, endometrium, stomach or kidney cancer, or tumour metastasis. This sequence represents an anitsense oligonucleotide used to modulate expression of urokinase plasminogen
                                                                                                                    The invention relates to a method of diagnosing juvenile polyposis (JP) or colorectal carcinoma. The method involves obtaining a sample from a subject and determining the loss or alteration of a functional BMPRIA gene in cells of the sample. The method is useful in diagnosing or treating JP or colorectal carcinoma. The invention is also useful in ge therapy. The present sequence is BMPRIA exon specific PCR primer, used illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                               Diagnosing or treating juvenile polyposis or colorectal carcinoma, comprises obtaining a tissue or fluid sample from a subject and determining the loss or alteration of a functional BMPRIA gene in
                                                                        Sequence
                                                                                                                                                                                                                                                                                                                         Example 1; Page 74; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                           determining the loss or alteration of the sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-MAY-2003
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                                                                        BP;
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0.2%;
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                                                                        3 C; 6
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3
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Pred.
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Pred. No. 2
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                                                                           T; 0 U; 0 Other;
14.8; DB 1;
No. 2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BMPR1A; gene therapy;
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                       Length
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PCR primer, used to
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ACC86760/c
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Best Local &
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                                                                                                                                                                                                                                    The invention relates to a method of diagnosing juvenile polyposis (JP) or colorectal carcinoma. The method involves obtaining a sample from a subject and determining the loss or alteration of a functional BMPRIA gene in cells of the sample. The method is useful in diagnosing or treating JP or colorectal carcinoma. The invention is also useful in gene therapy. The present sequence is a PCR primer used in BMPRIA exon mutation analysis. This primer is used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                      Diagnosing or treating juvenile polyposis or colorectal carcinoma, comprises obtaining a tissue or fluid sample from a subject and determining the loss or alteration of a functional BMPRIA gene in cells of the sample.
                                                                                                                                                                                                                                   mutation analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Juvenile polyposis;
diagnosis; PCR; pri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD53846 standard;
                            04-AUG-2003
                                                                                                                                                                                                      Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
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                                                                      ACC86760
                                                                                                                                                                                                                           invention
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                                                                                                                                                                                                      A.
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                                                                                                                                                                                                        0 Other;
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Human VEGFR-1 chimeric phosphorothioate

oligonucleotide

SEQ

ID NO:55

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The present invention describes a compound (C) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding vascular endothelial growth actor receptor-1 (VEGFR-1), where the compound inhibits the expression of VEGFR-1 and specifically hybridises with the nucleic acid encoding VEGFR-1 are viewed an annucleobase portion of an active site on the nucleic acid molecule encoding VEGFR-1. Also described: (1) a composition of acid molecule encoding VEGFR-1. Also described: (1) a composition of vEGFR-1 in cells or tissues by contacting the cells or tissues with (C) that the expression of VEGFR-1 is inhibited; and (3) treating an animal having a disease or condition associated with VEGFR-1 by administering (C) to the animal so that the expression of VEGFR-1 is inhibited; and can be used in antisense cytostatic and antisinflammatory activities, and can be used in antisense gene therapy. The antisense compounds are useful for modulating the expression of VEGFR-1 and for treating diseases or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 such as hyperprises or conditions associated with the expression of VEGFR-1 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             with the expression of VEGFR-1, such as hyperproliferative disorders (e.g. cancer), rheumatoid archiritis, or diseases or conditions involving angiogenesis. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. The present sequence represents a human VEGFR-2 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vascular endothelial growth factor receptor 1; VEGF receptor; VEGFR; inhibitor; cytostatic; anti-rheumatic; anti-archiritic; anti-anglogenic; anti-inflammatory; anti-sense gene therapy; hyperproliferative disorder; cancer; rheumatoid arthritis; anglogenesis; infection; inflammation; tumour formation; phosphorothioate; 2'-O-methoxyethyl; 2'-MOE; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotide targeted to a nucleic acid encoding vascular endothelial growth factor receptor-1, useful for diagnosing treating cancer, rheumatoid arthritis, or diseases or conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
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                                                                                                                                                                                                                                                                                                                            2 A;
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/note= "This oligonucleotide has a phosphorothioate backbone and 2'-0-methoxyethyl (2'-MOE) wings at the and 3' ends, which are 5 nucleotides in length. Also cytidine residues are 5-methylcytidines"
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                                                                                                                                                                                  The present invention relates to the sequencing of genomic DNA encoding human oestrogen receptor alpha (ESR1) protein. The gene encoding human ESR1 is located on chromosome 6. The invention provides the genomic structure of the ESR1 gene and novel single nucleotide polymorphisms (SNPB)/haplotypes in the genes. The polymorphisms/haplotypes can lead to a variety of disorders (such as cancer, osteoporosis, and cardiovascular disorders) that are mediated by a variant oestrogen receptor. The invention provides methods of detecting ESR1 polymorphisms/haplotypes in a sample, methods of determining a risk of having or developing a disorder mediated by a variant oestrogen receptor and methods for screening compounds useful for treating such disorders. ABX33756-ABX33782 represent sequencing primers for the human ESR1 gene
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22-FEB-2000;
20-OCT-2000;
24-JAN-2001;
13-MAR-2001;
05-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cardiovascular disorder; variant oestrogen receptor; ESR1 haplo ESR1 polymorphism detection; cytostatic; osteopathic; cardiant; sequencing; primer; ss.
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2000US-00692414.
2001US-00768184.
2001US-00804076.
2001US-00826314.
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eptor; ESR1 haplotype;
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                                                                                                                                The invention relates to antisense compounds, compositions and methods for modulating the expression of PCTAIRE protein kinase 1 (also known as PCTAIRE-1, PTCK1 and crk5). The antisense oligonucleotide is useful for treating an animal having a disease or condition associated with PCTAIRE protein kinase 1, particularly a hyperproliferative disease or a neurological disease. These diseases include thrombocytopaenia, mental retardation, wiskort-Aldrich syndrome, retinitis plymentosa, dystonia with Parkinsonism, sideroblastic anaemia, X-linked Charcot-Marie-Tooth disease, or incontinentia pigmenti. The antisense oligonucleotide is particularly useful for inhibiting the expression of PCTAIRE protein kinase 1 in cells or tissues. It is useful for diagnostics, prophylaxis, or as research reagents or kits. The present sequence is an antisense oligonucleotide targetted to human PCTAIRE protein kinase 1 DNA. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotides for modulating PCTAIRE protein kinase gene expression, particularly useful for treating hyperproliferative neurological disorders for example, mental retardation, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-DEC-2001; 2001US-00017621
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22-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                             05-APR-2002; 2002US-00116949
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                     single strand conformation polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human calcium channel alpha2delta SSCP primer c163E6R
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99US-00470443
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The invention relates to a substantially purified amino acid sequence comprising at least a portion of human calcium channel alpha2delta subunit splice isoform 1, splice isoform 2 sequence or splice isoform 3, or their variants, and their encoding nucleic acids (or their compound for modulating calcium channel activity, an antibody which binds to the calcium channel or its variants and producing a transgenic non-thuman animal (where the animal expresses a reduced level of calcium channel alpha 2delta subunit relative to a corresponding wild-type animal. The calcium channel proteins are useful for generating an animal (where the animal expresses a reduced level of calcium channel alpha 2delta subunit relative to a corresponding wild-type animal. The calcium channel proteins are useful for generating an comit of their portions or their portions). The transgenic animal (preferably a rodent e.g. mouse) is useful for identifying a therapeutic compound for treating a transgenic animal calcium channel proteins or their portions. The archival carcinoma or non-small calcium channel, breast cancer, naspoharyngeal cancer, cervical cancer, cancer, a neurological disease, especially epilepsy, a transgenic animal allowers, is a neurological disease, bultinfarct dementia, transpendent cancer, a neurological disease, wultinfarct dementia, Example New substantially pure human calcium channel alpha2delta subunit splice isoform 1, 2 and 3 sequence useful in preventing, treating and diagnosing cancer, neurological disorders and cardiovascular disease. 7; 25; English.

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Best Local
This invention relates to novel chimeric antisense oligonucleotides that specifically hybridise to and inhibit the expression of the nucleotide binding oligonucleotide domain 1, NODI protein. NODI, also known as CARDA (caspase associated recruitment domain 4) is a domain that is involved in the elimination of cells via programmed cell death and in the host defence against pathogens, i.e. it works to regulate apportosis. Apoptosis is a naturally occurring process, however, if it becomes overstimulated it can lead to cell loss and neurodegenerative conditions including Alzheimer's, Parkinson's, amyotrophic lateral sclerosis (ALS), retinitis pigmentosa and blood cell disorders. Conversely, insufficient apoptosis can contribute to the development of cancer, autoimmune disorders and viral infections. The present invention describes antisense oligonucleotides that can modulate NODI expression (and variants to thereof), such that these compounds, via gene therapy, can be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       arrhythmia, angina pectoris, hypoxic damage to the cardiovascular system, ischaemic damage to the cardiovascular system, myocardial infarction, and congestive heart failure, or Lambert-Eaton myasthenic syndrome. The proteins and nucleic acids are useful in the diagnosis, prevention and treatment of the above mentioned diseases. The human gene for the calcium channel is located on chromosome 3p21.3. The present sequence is an SSCP (single strand conformation polymorphism) primer used to detect polymorphisms in the calcium channel alpha2delta subunit gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compound, comprising a sequence targeted to a nucleic acid encoding nucleotide-binding oligomerization domain 1 (NOD1), useful for preparing a composition for treating hyperproliferative disease, e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; Page 75; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-DEC-2001; 2001US-00006883
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2002; 2002WO-US038606
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or disorder of nervous system which involves administering a modulator to condulate an activity of a neural stem cell or a neural progenitor cell in condulate an activity of a neural stem cell or a neural progenitor cell in condulate an activity of a neural stem cell or a neural progenitor cell in condulate an interaction between EphA7 and ephrin-A5 cor an interaction between EphA7 and ephrin-A2. The method is useful for alleviating a symptom of a disease or disorder of the nervous system, ce.g., drug and alcohol abuse, neurological traumas, or neurodegenerative, neural stem cell, neural progenitor, ischaemic, affective, neuropsychiatric or learning and memory disorders, such as Parkinson's disease, Huntington's disease, Alzheimer's disease, spinal ischaemia, amyotrophic lateral sclerosis, ischaemic stroke, spinal cord injury or cancer-related brain/spinal cord injury, schizophrenia, psychoses, componer control of the conductor of the nervous syndromes of disorders, spinal schaemia, stress and related syndromes, cognitive function disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide sequence is the chimeric antisense oligo used to inhigexpression of human NOD1, the aim of the invention. Note that it has terminal five nucleotide 2-methoxyethyl (2'-MOE) wings separated by ten deoxynucleotide gap. The oligonucleotide backbone is phosphorothi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EphA7; ephrin-A5; ephrin-A2; borderline personality disorder; ischaemia; epilepsy; trauma; infection; multiple sclerosis; autism; cerebral palsy; Huntington's disease; Alzheimer's disease; schizophrenia; gene therapy; memory disorder; Parkinson's disease; phobia; dementia; sleep disorder; amyotrophic lateral sclerosis; attention deficit disorder; depression;
                                                                                                                                                                                                                                                   The
                                                                                                                                                                                                                                                                                                                     Alleviating a symptom of a disease or disorder of the nervous administering a modulator of neural stem or neural progenitor activity in vivo to a patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-NOV-2001;
02-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-441543/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                 invention relates to a method for alleviating a symptom of a diseas
disorder of nervous system which involves administering a modulator
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human;
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The invention relates to a method for alleviating a symptom of a disease or disorder of nervous system which involves administering a modulator to modulate an activity of a neural stem cell or a neural progenitor cell in vivo to a patient suffering from the disease or disorder of the nervous system (the modulator disrupts an interaction between EphA7 and ephrin-A5 or an interaction between EphA7 and ephrin-A5 and interaction between EphA7 and ephrin-A5 or alleviating a symptom of a disease or disorder of the nervous system, e.g., drug and alcohol abuse, neurological traumas, or neurodegenerative,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Huntington's disease; Alzheimer's disease; schizophrenia; gene therapy; memory disorder; Parkinson's disease; phobia; dementia; sleep disorder; amyotrophic lateral sclerosis; attention deficit disorder; depression;
                                                                                                                                                                                                                                                                                                                                                                          Alleviating a symptom of a disease or disorder of the nervous administering a modulator of neural stem or neural progenitor activity in vivo to a patient.
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02-JUL-2002;
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                                                                                                                                                                                                                                              Diamond
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR;
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The invention relates to a composition comprising a sustaine chicken embryonic stem cells with a transgene stably integras genome of substantially all of the progeny of the embryonic The embryonic stem cell progeny contribute to a somatic tiss

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New compositions comprising sustained chicken embryonic stem (culture having a transgene integrated into genome of ES cell puseful in producing engineered chickens for avian transgenics,

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RESULT 3089
ADC37131
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Best Local
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The invention relates to a novel method for directly amplifying nucleic acids in a sample without having to isolate it. The method comprises a step to inhibit the breakdown of nucleic acids. The invention further comprises: a solution for treating the nucleic acid sample; a method for treating the sample; a reagent for detecting nucleic acids; a nucleic acid detection system; and a method for detecting cancer metastasis. This polynucleotide represents a PCR primer used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     into a chicken embryo. Compositions comprising cultures of embryonic chicken embryonic stem cells are useful in producing engineered chickens for avian transgenics, including protein production for pharmaceutical industry, production of chickens that deposit human antibodies in their eggs, and site-specific modification of the avian genome for other applications. The cultures may be used in the selection of desirable phenotypes in chimeric animals and modifications to the genome of chicken embryonic cells to introduce exogenous DNA into a chimeric offspring. Sequences ACF36177-186 represent primers used in a PCR analysis of transgenes for the presence of an unrearranged human heavy chain locus
                                                                                                                                                                                                                                              breakdown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-JAN-2002;
13-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-JAN-2003; 2003WO-JP000060.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CK19 forward primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC37131 standard;
                                                                                                                                                                                              Example 7; SEQ ID NO 20; 56pp; Japanese
                                                                                                                                                                                                                                                                    Directly amplifying nucleic acids
                                                                                                                                                                                                                                                                                                                      WPI; 2003-587284/55
                                                                                                                                                                                                                                                                                                                                                                 Yamagata
                                                                                                                                                                                                                                                                                                                                                                                            Tada S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003060116-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC37131;
                                                                                                                                                                                                                                                                                                                                                                                                                                            (SYSM-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric chicken
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplifying nucleic acid; inhibit; breakdown; cancer metastasis;
mer; ss; CK19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGAAAGGAGAGCACTGT 6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                            Yoshida T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGAGAGGAGAGCACAGT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002JP-00002046.
2002JP-00329958.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
                                                                                                                                                                                                                                                                                                                                                                                          Shohmi K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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Pred. No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                       Nishida M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                    in a sample
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                                                                                                                                                                                                                                                                                                                                                                                          Nakabayashi
                                                                                                                                                                                                                                                                    by inhibiting nucleic acid
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Query Match Best Local S Matches 16

Local Similarity mes 16; Conserv

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CC single nuclectide polymorphism (SNP) chosen from: (1) a nucleic acid of CS single nuclectide polymorphism (SNP) chosen from: (1) a nucleic acid of CC (1) or its complement under highly stringent hybridistions. (CC (1) or its complement under highly stringent hybridistions. (CC (1) or its complement under highly stringent hybridistion conditions. (CC (1) contiguous nuclectides of a nuclectide sequence of S. salar SNPs, O. filoticus SNPs, O. inloticus microsatellites, Atlantic halibut SNPs, Cod (1) polymorphic sites and seabass polymorphic sites, or their complement; (2) CC a primer pair (III) suitable for use in PCR, comprising two (II) capable of anymorphic sites and seabass polymorphic sites, Atlantic halibut SNPs, cod (1) comprising a nuclectide sequence chosen from S. salar SNPs and, O. (1) capable (1) comprising a nuclectide sequence chosen from S. salar SNPs and, O. (1) contiducts sNPs, O. niloticus microsatellites, Atlantic halibut SNPs, cod (2) corigin of fish sample comprising providing a parentage genotype database (2) corigin of fish sample comprising providing a parentage genotype database (2) candidate parent genotype represents a distinct origin, and comparing a comprising a collection of the candidate parent genotype identifies (2) the sample genotype and one of the candidate parent genotype identifies (2) to the origin of the sample. (M1) is useful for detecting nucleic acid molecule comprising sucleic acids with one or more (2) contident from nucleotide sequence of S. salar SNPs and O. niloticus (2) contident for detecting nucleic acid that hybridises to (II) (II) is contident for detecting nucleic acid molecule comprising a polymorphic contacting contacting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 3090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ... 12. 989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes an isolated nucleic acid (I) comprising single nucleotide polymorphism (SNP) chosen from: (1) a nucleic acid of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated nucleic acid molecule comprising single nucleotide polymorphism associated with fish, useful for forming PCR primers which are used for detecting single nucleotide polymorphisms in fish nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lie O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Orechromis niloticus; Atlantic halibut; microsatellite; polymorphic site; seabass; salmonidae; Tilapia; rainbow
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-627388/59
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16-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JAN-2003; 2003WO-IB000112
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2002US-0404200P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Salmo salar;
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RESULT 3091
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ID ADB0223
AC AAD6223
AC AAD622
AC AC AAD62
AC AC AAD62
AC AC AAD62
AC AC AAD62
AC AAAC6
AC AAC6
AC ACC6
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acids with one or more (II) which is derived from O. niloticus microsatellite, O. niloticus SNPs, Atlantic halibut SNPs, cod polymorphic sites or seabass polymorphic sites, and identifying a nucleic acid that hybridises to (II) (III) is useful for detecting nucleic acid molecule comprising a microsatellite sequence in sample. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAD62234 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                   New antisense oligonucleotides targeted to nucleic acids encoding hematopoietic cell protein tyrosine kinase, useful for diagnosing treating cancer (e.g. leukemia), inflammation, diabetes or viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2003125275-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Haematopoietic cell; tyrosine kinase; hyperproliferative disorder;
cancer; therapy; inflammation; diabetes; viral infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD62234;
  The invention relates to a compound targetted to a nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                  04-DEC-2001; 2001US-00007010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     phosphorothioate
                                                  Example 15; Page 27;
                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7021 ACAGAGGAAAATAGGAAA 7038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapy; inflammation; diabetes; viral ir cytostatic; virucide; antisense therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                       2001US-00007010
                                                                                                                                                                                                                                                                                                                       PHARM INC.
                                                                                                                                                                                                                                                                       Dobie KW;
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/*tag=
/mod_bas
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/note= "Phosphorothioate backbone;
methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       backbone;
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                                               59pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "2'-O-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "2'-O-methoxyethyl (2'-MOE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense;
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              encoding haematopoietic cell protein tyrosine kinase. The compound inhibits the expression of haematopoietic cell protein tyrosine kinase and it specifically hybridises with the nucleic acid molecule encoding the tyrosine kinase or with at least an 8-nucleobase portion of an active site on the nucleic acid molecule encoding the tyrosine kinase. The antisense compounds are useful for modulating the expression of haematopoietic cell protein tyrosine kinase and treating diseases or conditions associated with the expression of the tyrosine kinase, such as hyperproliferative disorders (e.g. cancer), inflammation, diabetes or a viral infection. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological
                                                                                                                                         Investigating male genetic infertility, useful for diagnosis e.g. for assessing suitabilty for in vitro fertilization, based on multifactorial analysis of infertility-related mutations.
                                                                                                                                                                                                                                                                                                                                                                                                                                       cystic fibrosis tra
Kallmann syndrome;
CYP21; microarray;
                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 7 A; 3 C; 4 G; 6 T; 0 U;
                                                                                                                                                                                                                             Cullen
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                                                                                                                                                                                                                                                                                                                                                                                                                           oligospermia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                azoospermia factor; Y-chromosome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer; male infertility; infertility-associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD42189;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        kinase antisense oligonucleotide
                                                                                                               Claim 13; SEQ ID NO 50; 110pp; German
                                                                                                                                                                                                                                                                                  10-DEC-2001; 2001DE-01060563
                                                                                                                                                                                                                                                                                                                                           19-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                     WO2003050299-A2
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human infertility associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD42189 standard;
                                                                                                                                                                                                                                                         (OGHA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7140 CCAGCCTAATTGGTATGT 7157
                                                                                                                                                                                                  2003-505402/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                           ۵,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fibrosis transmembrane conductance regulator; CTFR;
                                                                                                                                                                                                                                                         OGHAM GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCAGCCTAATATGTATGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                          quantitative
                                                                                                                                                                                                                                                                                                                                                                                                                                       KALI; androgen resistance; steroid 21-hydroxylase; quantitative trait locus; in vitro fertilization;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mutation;
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This invention describes a novel method for investigating genetic infertility or predisposition in males. The method involves selecting least two infertility-associated mutations which are recessive or intermediate that are associated with infertility in the heterozygous state and/or only in the homozygous state. Preferably at least one azoospermia factor is detected which may be lost by microdeletions in intervals 5 or 6 of the Y-chromosome. Also any of several hundred

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RESULT 3093
ADD81662
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CC conductance regulator (CTFR), Kallmann syndrome (KAL1), androgen
CC resistance (AR) or steroid 21-hydroxylase (CYP21) genes may be detected.
CC Probes for the mutated genes and/or native nucleic acid, or their
CC complementary strands, are fixed to a carrier, particularly as a
CC microarray, then tested for hybridization with oligonuclectides from or
CC synthesized from, a patient sample and hybridization detected.
CC Multifactorial analysis is by standard statistical methods, particularly
CC the quantitative trait locus method. The method is used to diagnose
CC conjunction with in vitro fertilization programs, e.g. for assessing
CC conjunction with oligospermia for possible application of the
CC conjunction with oligospermia for possible application of the
CC intracytoplasmic sperm injection method. Analysis of many mutations
CC subjects with oligospermia for possible application of the
CC intracytoplasmic sperm injection method. Analysis of many mutations
CC conjunctions). A chip for analyzing genetic infertility in males comprises
CC oligonucleotides that represent known mutations (nonsense or missence,
CC insertions, allelic variants deletions or rearrangements) in the cystic
CC fibrosis transmembrane conductance regulator. Kallmann syndrome. androgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 16;
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Best Local
Predicting potential of oligonucleotides to hybridize to target nucleotide sequence comprises determining and evaluating for each oligonucleotide a parameter predictive of the oligonucleotides abihybridize with target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fibrosis transmembrane conductance regulator, Kallmann syndrome, androgen resistance and steroid 21-hydroxylase genes. ADD42140-ADD42633 represent oligonucleotides used in the microarray described in the method of the invention. NOTE: there are no SEQ ID's 133, 472 or 473 represented in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003054346-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV PRT antisense derived probe #591.
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                                                                                                                                                                                                                                                                                                                                                                             (SHAN/)
                                                                                                                                                                                                                                                                                                                             (DELE/)
                                                                                                                                                                                                                                                                                                                                                     (MOLB/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide hybridisation potential; efficient hybridisation; ge array; minimum oligonucleotide synthesis; probe.
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WOLBER P K.
DELENSTARR G (
WEBB P G.
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                                                                                                                                                                                                                                                             KINCAID R H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               virus
                                                                                                                                                                                                          Delenstarr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8;
Pred. No. 26
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                                                                                                                                                                                                    GC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2e+03;
                                                                                                                                                                                                       Webb
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                              ability
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RESULT 3094
ADE39777/c
ID ADE3977
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Best Local (
assaying a cellular material sample from known origin. It can be used to determine if an animal by susceptibility to porcine (especially pigs) to pRRSV and to compare susceptibility to a known origin. It can be used to determine the susceptibility as frown susceptibility to a known susceptibility to a known susceptibility to a known susceptibility as the susceptibility to a known origin. It can be used to determine if an animal of the same origin. It can be used to determine if an animal of the same origin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of predicting the potential of oligonucleotides to hybridise to target nucleotide sequences. The method is useful for predicting the potential of an oligonucleotide to hybridise to a target nucleotide sequence, e.g. RNA or DNA or a sequence that contains chemically modified nucleotides. The method is also useful for predicting the potential of the oligonucleotides to hybridise to a complementary target nucleotide sequence. The method is useful to predict efficient hybridisation oligonucleotides for each of multiple target sequences therefore very large arrays may be constructed and tested with minimum synthesis of oligonucleotides. The present sequence represents a HIV PRT antisense derived probe.
                                                                                                                                                                                                                                                                                                                                                                               Determination of susceptibility to porcine reproductive and respiratory syndrome virus non-invasively useful e.g. to breed pigs with low susceptibility or classify infection resistance in an animal, by assaying the control of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             porcine reproductive and respiratory syndrome virus; PRRSV; susceptibility; CD 151; susceptibility standard; PRRSV infection; vaccine; vaccine virus stock; non-simian vaccine; xenotransplantion; non-simian cell line; drug testing; transformed cell line; porcine; pig;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-811729/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kapil S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2001; 2001US-00772044.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE39777;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADE39777 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 735; 423pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KAPI/) KAPIL S.
(SHAN/) SHANMUKHAPPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gg.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred.
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RESULT 3095
AD240361
ID AD24036
AC AD24036
CW ANTIGER
CW ANTIGER
CW ANTIGER
CW ANTIGER
CW ANTIGER
CW ASTHMAR
CW ASTHMAR
CW ASTHMAR
CW AD20030
CW AUG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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Best Local S
Matches 16
02-AUG-2001;

03-AUG-2001;

07-AUG-2001;

08-AUG-2001;

09-AUG-2001;

13-AUG-2001;

17-AUG-2001;

17-AUG-2001;

20-AUG-2001;

20-AUG-2001;

21-AUG-2001;

21-AUG-2001;

23-AUG-2001;

23-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NOVX; cardiant; antiarteriosclerotic; hypotensive; cytostatic; anorectic; antidiabetic; immunosuppressive; anti-HTV; neuroprotective; nootropic; antiparkinsonian; antiasthmatic; gynaecological; cardiomyopathy; atherosclerosis; hypertension; cancer; obesity; diabetes; AIDS; multiple sclerosis; graft-versus-host disease; Alzheimer's; Parkinson's;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reverse Ag4809 RT-PCR primer used to amplify human NOV RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE40361 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               asthma; fertility disorder; vaccine; gene therapy; tissue typing; human; NOV; ss; primer; PCR; RT-PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                            02-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-AUG-2003
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2001US-0309501P.
2001US-0310549P.
2001US-0310549P.
2001US-0311992P.
2001US-0311992P.
2001US-0311992P.
2001US-0313415P.
2001US-0313415P.
2001US-0313415P.
2001US-0313415P.
2001US-0313415P.
2001US-0313415P.
2001US-0313415P.
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Pred. No. 2e+03;
Nismatches
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TXXXXFFXXXXXXX

HBV target 25-MAR-2003 26-MAY-1994

sequence

10 entry)

(revised) (first en

RNA; enzyme; enzymatic RNA molecule; ERM; cleave; RNA; mRNA; H picornavirus; HIV; immunodeficiency virus; hepatitis B virus; papilloma virus; HPV; Epstein-Barr virus; EBV; TCLV; T-cell leukaemia virus; hepatitis C virus; HCV; cytomegaloviru

HCV; cytomegalovirus;

밁 S

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18

2337 CCATCACACCCGCCTTTT 2354

AAQ50784 standard;

RNA;

21 BP

AAQ50784;

Matches Best

16;

Conservative

0

Mismatches

2

<u>,,</u>

Gaps

0

Local

Similarity

88.9%;

Pred.

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2e+03;

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Kekuda R, Zerm. Shimkets RA, Zerm. Shimkets RA, Zerm. I Voss EZ, Boldog FL, Gorman I, Was Zanger SR, Ellerman K, Ma Tainger SR, Lepley DM, I
Query Match
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17-SEP-2001; 2001US-0322716P.
21-SEP-2001; 2001US-0323994P.
14-DEC-2001; 2001US-034233P.
05-PEB-2002; 2002US-0354591P.
19-MAR-2002; 2002US-0355478P.
19-APR-2002; 2002US-0373814P.
19-APR-2002; 2002US-0373825P.
19-APR-2002; 2002US-0373825P.
23-APR-2002; 2002US-0373825P.
23-APR-2002; 2002US-0374632P.
07-UUN-2002; 2002US-0386971P.
01-AUG-2002; 2002US-03210172.
                                                                              The invention relates to a novel NOVX polypeptide. The polypeptide of the invention demonstrates cardiant, antiarteriosclerotic, hypotensive, cycostatic, anorectic, antidiabetic, immunosuppressive, anti-HIV, neuroprotective, noctropic, antiparkinsonian, antiathmatic and gynaecological activities and may be useful in diagnosing, treating or preventing NOVX-associated disorders including cardiomyopathy, atherosclerosis, hypertension, cancer, obesity, diabetes, AIDS, multiple sclerosis, graft-versus-host disease, Alzheimer's disease, Parkinson's disease, asthma or fertility disorders. Furthermore, the polypeptides may be utilised as vaccines whilst the nucleic acids may be used as hybridisation probes, in gene therapy, chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that of the RT-PCR primer of the invention which was used to amplify human NOV man
                                                                                                                                                                                                                                                                                                                                                                                         New NOVX polypeptides and nucleic acids, useful for preventing treating NOVX-associated disorders, e.g. cancer, cardiomyopathy atherosclerosis or diabetes, and in chromosome mapping, tissue to
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-663472/62.
                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                     Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            r CE, Patturajan M, Pena CEA, Rieger DK;
rhusen BD, Li L, Ji W, Padigaru M, Casman SJ;
FL, Gorman L, Leite MW, Vernet CAM, Anderson i
Gerlach VL, Hjalt T, Rastelli L, Spytek KA;
erman K, Malyankar UM, Macdougall JR, Stone DJ;
epley DM, Burgess CE, Majumder K, Wolenc AR;
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                                                                                                                                                                                                                                                                                                                                     267; 560pp;
 0.2%;
                                    C; 1
                                    G; 5 T;
 Score
                                                                                                                                                                                                                                                                                                                                     English
 14.8;
                                    0 U;
                                      0 Other;
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   Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                          typing
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Query Match
Best Local S
Matches 6
                                                                                                                                               The sequences (AAQ50775-99) and (AAQ52551-8) are pref. hepatitis B virus target sequences for enzymatic RNA molecules. The RNA molecules are complementary to a substrate binding region in the specified gene target. They also have enzymatic activity, in that they specifically cleave RNA in the target. The ERMs interfere with viral replication and therefore have anti-viral properties. They can be used to attenuate viruses to be used in vaccines. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                   WPI; 1993-386599/48.
                                                                                                                   Sequence
                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                            Mamone
                                                                                                                                                                                                                                                                                                                                                       Draper
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          influenza virus; HSV; herpes simplex virus; vector; immune response;
antibody; ribozyme; viral RNA; treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                              (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-1992;
14-MAY-1992;
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14-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAY-1992
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14-MAY-1992;
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14-MAY-1992;
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14-MAY-1992;
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                                                                                                                                         25-MAR-2003
field.)
                                                                                                                                                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                                                           JA,
                        N
                                                                     Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                              RIBOZYME
                                              TGCCTTGTTTCCTTCTAT 2921
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                        uGACUUCUUUCCUUCUAU 19
                                                                                                                                                                                                                                                                                expression.
                                                                                                                                                                                                                                                                                           RNA molecules -
                                                                                                                   BP;
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                                                                       Conservative
                                                                                                                                                                                                                                                          8; 287pp; English.
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3-00963322.
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3-00936086.
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3-00923738.
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-00882922.
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-00884333.
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-00884073.
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                                                                     Score 14.8; D
Pred. No. 2.1e
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                                                                                                                                                                                                                                                                                           inhibit viral replication,
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                                                                                 2.1e+03
                                                                                           DB 1;
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AAQ65789
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                                                                                                                                                                                    RESULT 3098
AAQ55255/c
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Best Local S
Matches 16
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20-DEC-1994
                                           Ribosome; grapevine; Vitis; IGS region; rDNA; polymorphism; grape cultivar; probe; primer; detection; primer; amplify; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                   gene
WO9401580-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting genetic pre-disposition to osteoarthritis - and other diseases involving mutation in cartilage protein genes, by amplification and analysis of DNA and comparison with standards.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ65789 standard;
                                                                                           Grapevine
                                                                                                                 25-MAR-2003
14-JUL-1994
                                                                                                                                                   AAQ55255;
                                                                                                                                                                         AAQ55255 standard; DNA;
                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                  Claim 18 claims primers for use in detecting mutations in a mammalian gene for a structural protein of cartilage comprising a sequence identified in Table I (Page 18-31). Table I includes 179 primer sequences (see AAQ65728-Q65906). The following details are given for primer 55:
Alt. code: DH-47 Region/exon: 21 Direction: antisense Primer position: 9660 (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 18; Page 23; 112pp; English.
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                                                                                           rDNA detection
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Ahmad NN;
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(first entry)
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(first entry)
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reaction; PCR; osteoarthritis; cartilage;
                                                                                                                                                                                                                                                                                                                            A;
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                                                                                                                                                                                                                                                                                                                            G;
                                                                                          primer #5.
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Pred. No. 2.1e
0; Mismatches
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Best Local S
Matches 16
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  Oligo: ribonucleotide analogue for identifying new antibiotics -
                                                    WPI; 1996-151324/15:
                                                                                                                                                                                                       23-AUG-1994;
05-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analogue; ligand binding; 16S rRNA; parental molecule; conformation; stem-loop structure; stable; helix; nucleotide clamp; decoding region; mRNA; tRNA; A site; ribosome; protein synthesis; assay; antibiotics;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E.coli tRNA(Pro) anticodon stem-loop used for generating analogues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-OCT-1996
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                                                                                                                                                                                                                                                                                   23-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                       WO9606106-A1
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16; Conserv
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95US-00498402.
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Pred. No. 2
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  are generated from two sections, the first section corresp. to sequences in the parental RNA mol., whilst the second section consists of an artificial nucleotide structure (i.e. not found in the parental RNA) that can combine with the first section to stabilise the analogue. The second structure also contains a stem-loop structure and a second nucleic structure able to form a stable base-paired helix, designated the nucleotide clamp. The analogues pref. contain the complete sequence of the 16S rRNA decoding region (the decoding region being the region which correctly aligns the mRNA with the tRNA in the "A site" of the ribosome during protein synthesis). The sequences ATI6748-52 were used to generate analogues of the invention esp. for aminglycoside protection assays. The analogues can be used in assays to detect potential ligands e.g. new antibiotics or anti-viral agents, that could bind to the parent
                                                                                                                                                                                                                                                                                                                                                                     polycystic kidney disease; PKD1; autosomal dominant; ADPKD; mutation; exon; short arm; chromosome 16; regated region; alternative splicing extracellular matrix proteins; antibody; detection; diagnosis; mini gene therapy; single strand conformational polymorphism analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             binding pattern as parental RNA derivative, therapeutic activity of cpd.
                                                    DNA encoding poly-cystic therapy of ADPKD, and in
                                                                                                                                                                                                                                                                     21-DEC-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT11723;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel analogues able to assume the same conformation as the parent mol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 18; 57pp; English.
                         Example; Page 68; 126pp; English.
                                                                                                                             Reeders
                                                                                                                                                                                                  30-MAR-1995;
                                                                                                                                                                                                               03-JUN-1994;
                                                                                                                                                                                                                                           02-JUN-1995;
                                                                                                                                                                                                                                                                                                   W09534573-A1
                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                         SSCP;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Polycystic kidney disease 1 gene antisense primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2003
18-JUL-1996
                                                                                               WPI; 1996-049618/05
                                                                                                                                                       (MILL-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT11723 standard;
                                                                                                                                                                      (BGHM )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4187 GGTTATCGCCCCAAGATG 4204
                                                                                                                                                                                                                                                                                                                                                        primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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                                                                                                                           s
                                                                                                                                                       BRIGHAM & WOMENS HOSPITAL.
MILLENIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                           Schneider M,
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(first entry)
                                                                                                                                                                                                94US-00253524
95US-00413580
                                                                                                                                                                                                                                           95WO-US007079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 4 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                      kidney disease
the evaluation
                                                                                                                           Glucksmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1;
Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 U;
                                                      gene product of treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and assays for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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    for pkD.

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The sequences given in AAT11709-29 are primers which were used in single-

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RESULT 3101
AAT86043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 16; Conserv
The sequences given in AAT86037-43 are primers which were used in the amplification of regions of the murine V-alpha-1 intron and the V-alpha-1 'promoter. The amplified sequences were used in the isolation of the antisense strand of the T-cell receptor gene which encodes a signalling protein which comprises a proline rich region, a YXXI/I motif and a TANYSNI motif. Expression of the signalling protein sequence is activated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense; T-cell receptor; signalling protein; intron; murine; V-alpha-1 gene; cytoplasmic tail; Ig-beta; P8.6; primer; amplif
                                                                                                                                                                                                                                                                    New T-cell receptor V-alpha gene signalling protein - encoded antisense strand of the V-alpha-1 gene, includes proline-rich
                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-132579/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                                                                                                                              Example 1;
                                                                                                                                                                                                                                               Tyr-X-X-Ile/Leu signal motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-FEB-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                          Maeda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amplifies
                                                                                                                                                                                           Page 12; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          murine V-alpha-1 intron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.9%;
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                                                                                                                                                                                              English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2887
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig-beta; P8.6; primer; amplify;
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RESULT 3103 AAV20817/c ID AAV2081

AAV20817

standard; DNA;

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AAV62909/c
                                     Query Match
Best Local Similarity
Matches 16; Conserv
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                                                                                                AAV62907-V62927 are PCR primers used in the amplification of a human galactokinase. This protein is used in a method to detect galactokinase mutations. This protein and its encoding nucleic be used in methods allowing the detection, diagnosis and treatm human galactokinase deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              murine V-alpha-1 gene. The encoded protein has a molecular weight of the cycles and has a high degree of homology (e.g. 70%) to the cytoplasmic for Ig-beta. The murine signalling protein P8.6 and its homologues, from human lymphoid cells, can be used to further elucidate the rol the T cell receptor V-alpha-1 gene locus in the functioning of T ce
                                                                             Sequence 21
                                                                                                                                                               Example
                                                                                                                                                                                               Detection
                                                                                                                                                                                                                  WPI; 1998-609232/51.
                                                                                                                                                                                                                                      Bergsma
                                                                                                                                                                                                                                                                               26-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                             deficiency:
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                                                                                                                                                                                                                                                                                                                                                                                                                          Human galactokinase
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                                                                                                                                                                                                                                                           (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                                                                                                                                              1; Col 35-36; 31pp; English.
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CAGGTCGCGGAGCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCTCCTTCCTCTTCACTG
                                                                                                                                                                                 of galactokinase mutations - based on comparison with wild-type ence or altered galactokinase activity.
                                                                               BP;
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                                      Conservative
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                                                                             3 A;
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                                                                              9 C; 7 G;
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he role of
f T cells
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                                                                                                                                                                                                                                                           Query Match
Best Local
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24-MAR-1997;
07-JUL-1997;
                                                                                                                                                                                                                                                                                                                                   This sequence is a primer for DNA encoding the human haematopoietic stem cell growth factor (SCGF) of the invention. The polypeptide of the invention is of mammalian origin and has haematopoietic stem cell growth factor SCGF activity, including burst-promoting activity (BPA) and granulocyte macrophage colony stimulating activity (GPA). The products can be used for treatment, diagnosis and analysis of haematopoietic cell disorders and bone marrow inhibition, e.g. by cytotoxic anticancer agents such as 5-fluorouracil. The products can also be used for amplification of haematopoietic cells in vitro, e.g. for use in marrow grafting and gene therapy by insertion of SCGF gene using a suitable therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Haematopoietic stem cell growth diagnosis of haematopoietic cell inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Haematopoietic stem cell growth factor; SCGF; b BPA; granulocyte macrophage colony stimulating GPA; haematopoietic cell disorder; bone marrow
 Polymorphism; human; inhibitor; cancer; treatment; cell growth; cell viability; loss of heterozygosity; precancerous condition; allele specific inhibitor; somatic cell; diagnosis; prevention;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV20817;
                                                                               30-NOV-1999
                                                                                                                                AAZ26119
                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 21; Page 49; 85pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-179383/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KYOW ) KYOWA
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                                                                                                                                                                                                21
                                                    polymorphic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Þ
                                                                                                                                                                                                                                                             Similarity
                                                                                                                                standard;
                                                                                                                                                                                                                         TGCCTGCATCAAACTGGA
                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human haematopoietic
                                                                                                                                                                                                                                                                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sugimura
                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               накко косуо кк.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96JP-00262252.
97JP-00087242.
97WO-JP002349.
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                                                                                                                                                                                                                                                                                                   σ
                                                    region 308.
                                                                                                                                                                                                                                                                                                   Ą
                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Þ
                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                                                                                                  5 C; 6
                                                                                                                                  21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      M.io
                                                                                                                                                                                                 4
                                                                                                                                                                                                                        2291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       with factor - useful for, e.g. treatment cell abnormalities and bone marrow
                                                                                                                                                                                                                                                                                                   G; 4 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ξ
                                                                                                                                                                                                                                                            Score 14.8;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8tem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell growth
                                                                                                                                                                                                                                                                                                  ₽;
                                                                                                                                                                                                                                                            2.1e+03
                                                                                                                                                                                                                                                                                                   0 Other
                                                                                                                                                                                                                                                                         DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     burst-promoting activity; g activity; gene therapy; w inhibition; human;
                                                                                                                                                                                                                                                   2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           factor
                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                 Gaps
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atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
WO9841648-A2
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Homo sapiens

98WO-US005419

97US-0041057P

(VARI-) VARIAGENICS

Housman D, Ledley FD, Stanton ď,

Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.

Disclosure; Fig 7; 605pp; English.

This invention describes a novel method for identifying an inhibitor CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient a first allele Specific inhibitor (ASI) targeted to an allele of a first essential gene CC specific inhibitor (ASI) targeted to an allele of a first essential gene CC calls of the patient are heterozygous for the first gene, the inhibitor CC calls of the patient are heterozygous for the first gene, the inhibitor CC calls of the patient are heterozygous for the first gene, the inhibitor CC calls of the galation and targets only one allelic forms of the gene CC used in the diagnosis, prevention and treatment of LOH disorders, e.g. CC cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic CC lesions, benign tumours, endometriosis, polycystic kidney disease, and CC graft versus host disease the mathod can also be used to remove CC malignant cells from bone marrow transplants. AAZ25812-Z26825 represent CC human polymorphic sites described in the method of the invention

Sequence 21 BP; 6 A; 4 C; 9 ဂ္ 2 T; 0 u; 0 Other;

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Matches
                        Query Match
Best Local
4198 CAAGATGGGGTCCAGGCT 4215
                         Similarity
                 Conservative
                           88.98;
                                  0.2%;
                 0
                           Pred.
                                  Score
                  Mismatches
                        14.8; DB 1;
No. 2.1e+03;
                                   DB 1;
                  Indels
                  <u>..</u>
                  Gaps
                  0
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RESULT 3105 밁 S \vdash CAAGAAGGGGACCAGGCT 18

30-NOV-1999 (first entry)

AAZ26693 standard;

DNA; 21

ВÞ

Human polymorphic region 882.

Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss. treatment; cell growth; precancerous condition;

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RESULT 3106
AAX35653
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for identifying an inhibitor CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC used for preventing the development of cancer in a patient having a CC used for preventing the development of cancer in a patient having a CC precancerous condition, by administraing to the patient a first allele CC specific lihibitor (ASI) targeted to an allele of a first essential gene CC present in cells of the precancerous condition, where the normal somatic CC cells of the patient are heterozygous for the first gene, the inhibitor CC is active on at least one but less than all allelic forms of the gene CC present in a population and targets only one allelic form present in the normal somatic cells, and the first gene. The products and methods can be CC used in the diagnosis, prevention and treatment of LOH disorders, e.g. CC cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic CC concers, benign tumours, endometriosis, polycystic kidney disease, and CC graft versus host disease. The method can also be used to remove considered the product of the product of the product of the premove transplants. AAZ25812-Z26825 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
W09911798-A1
                                                                                                                               Heparanase; hpa; modulator; heparin-binding growth factor; cellular response; cytokine; cell interaction; plasma lipocellular susceptibility; infection; disintegration;
                                                                                                                                                                                                                                             09-JUL-1999
                                                                                                                                                                                                                                                                                                            AAX35653 standard; DNA;
                                                                                              neurodegenerative plaque; wound healing; angiogenesis; restenosis; athersclerosis; inflammation; neurodegenerative disease; neutralis
                                                                                                                                                                                                    PCR primer used to amplify human heparanase cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 7; 605pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-521232/44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-SEP-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                   asma heparin; micrometastasis;
                                                                 primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                       5420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymorphic sites described in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                                                                                                                                                                                                                                                                      ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 BP; 10 A; 4 C; 5 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                      AAAAGGAAGTGAATCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAGCAAGAGAATCAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0041057P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US005419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                21
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                                                                                                                                                                                                                                                                                                                                                                                                                                     5437
                                                                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.8; DB 1;
Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                  autoimmune lesion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                   plasma lipoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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AAX18314
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       activity. The recombinant protein is used as a modulator of heparin-
binding growth factors, cellular responses to heparin-binding growth
factors and cytokines, cell interaction with plasma lipoproteins,
cellular susceptibility to viral, protozoal and bacterial infections or
disintegration of neurodegenerative plaques. Heparanase may be useful fo
conditions such as wound healing, angiogenesis, restenosis, and viral
infections. Mammalian heparanase can be used to neutralize plasma
heparin, and anti-heparanase antibodies may be applied for
immunodetection and diagnosis of micrometastases, and body fluids. The
present PCR primer was used to amplify hpa cDNA, in the course of the
                                                                                                                                               neuroblastoma; breast carcinoma; colon carcinoma; lymphoma; osteosarc smooth muscle cell hyperplasia; stem cell proliferation; Wilm's tumou stem cell differentiation; organ regeneration; organ differentiation;
                        01-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (INSI-)
(HADA-)
(FRIE/)
                                                  14-JAN-1999.
                                                                          WO9901560-A1
                                                                                                    Homo
                                                                                                                 Synthetic.
                                                                                                                                                                                         Telomerase; human; cancer; diagnosis; melanoma; skin cancer; leukaemia;
                                                                                                                                                                                                                   PCR primer for
                                                                                                                                                                                                                                               26-JUL-1999
                                                                                                                                                                                                                                                                         AAX18314;
                                                                                                                                                                                                                                                                                                 AAX18314 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes a polypeptide having heparanase (hpa) activity. The recombinant protein is used as a modulator of heparin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-SEP-1997;
02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-AUG-1998;
                                                                                                                                        PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-302255/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation
                                                                                                                                                                                                                                                                                                                                                                                          7413 CAGCAGCAGCAGCAGCAG 7430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human polynucleotide useful for treating angiogenesis, restenosis,
                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                   16;
                                                                                                                                                                                                                                                                                                                                                                 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7; Page 30;
                                                                                                                                                                                                                                                                                                                                                                 CAGGAGCAGCATCAG
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                                                                                                                                          88.
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A; 8 C; 6 G; 1 T; 0 U;
                                                                                                                                                                                                                                               (first entry)
                        98WO-US013835
                                                                                                                                                                                                                    telomerase coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-00922170.
98US-00109386.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98WO-US017954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Feinstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                 ₽₽
                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            & DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                  2.1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                    2
                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                               tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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01-JUL-1997;

97US-0051410P

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RESULT 3108
AAX79141/c
ID AAX7914
XX AAX7914
XX AAX7914
XX IT-AUG-
XX MSH6; M
KW Charact
KW Charact
KW Eimple
XX Synthet
OS Synthet
OS Arabido
XX MO99194
XX X IT-AUG-
XX ON-OST-
PF 09-OCT-
PR 10-OCT-
XX PA (RHON )
XX DOUTTIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-JUL-1997;
21-JUL-1997;
04-AUG-1997;
09-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence is a PCR primer for DNA encoding a truncated human telomerase of the invention. Primers that amplify the telomerase coding sequence can be used in a method for diagnosing cancer in a patient. The telomerase can be used for detection, diagnosis and drug screening. Inhibitors of telomerase activity can be used to treat cancers such as melanomas, other skin cancers, neuroblastomas, breast carcinomas, colon carcinomas, leukaemias, lymphomas, osteosarcomas or smooth muscle cell hyperplasias or skin growths. Enhancers of telomerase may be used to stimulate stem cell proliferation and differentiation (expansion of haematopoietic stem cells could be administered in the bone marrow transplant context). As well, many tissues have stem cells. Proliferation of these cells may be useful in wound healing, hair growth, treatment of disease such as Wilm's tumour, organ regeneration or differentiation after injury or diseases, nerve cell or brain cell growth following
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated vertebrate telomerase genes - used to develop treating cancers or for organ regeneration, nerve cell or b growth following injury or bone marrow transplantation.
                                                                                                                                                                                                                                                                                                                                                                         AAX79141 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-106060/09
                                                                                                                                                                                                                              characteristic; simple sequence
                                                                                                                                                                                                                                                                                                                   17-AUG-1999
                Doutriaux M,
                                                                                                  09-OCT-1998;
                                                                                                                                                           WO9919492-A2
                                                                                                                                                                                     Arabidopsis thaliana
                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                     Primer NGA1107-L for A.thaliana SSLP marker.
                                                                      10-OCT-1997;
                                                                                                                                                                                                                                                           MSH6; MutS homologue; plant; DNA mismatch
                                          (RHON ) RHONE-POULENC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 42; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bowtell D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A;
                                                                                                                                                                                                                                                                                                                   (first entry)
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97US-0053329P.
97US-0054642P.
97US-0058287P.
                Betzner AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BIOSYSTEMS
                                                                        97AU-00009745.
                                                                                                    98WO-EP006977.
                                                                                                                                                                                                                      microsatellite; primer; PCR; amplification; SSLP; 88; length polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
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                                            AGROCHIMIE.
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                                                                                                                                                                                                                                                                                                                                                                           21 BP
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                Freyssinet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DI
Pred. No. 2.1e.
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 T; 0
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                ຸດ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
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                 Perez
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 21;
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RESULT 3109
AAZ28937/c
ID AAZ2893
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AAX79066) and MSH6 (AAX79067) genes. These genes are Muts homologues (MSH) from plants and are involved in DNA mismatch repair. The DNA sequence can be used in processes for at least partially inactivating a DNA mismatch repair system of a plant, for increasing genetic variation in a plant, and for obtaining a plant with a desired characteristic primers AAX79105-X79160 represent 28 primer pairs used to amplify short allelic repeat fragments designated Simple Sequence Length Polymorphisms (SSLP). These fragments can be used as markers in the analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Reverse primer cmar8; paraplegin; human; hereditary spastic parap
HSP; mutation; diagnosis; treatment; neurodegenerative condition;
Amyotrophic Lateral Sclerosis; ALS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             system of
                                                                                                                                                                                                                                                                             08-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reverse primer cmar8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ28937;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ28937 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to the isolation of the Arabidopsis thaliana MSH3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 28; 117pp; English.
                                                                                                                                                                                                                   Ballabio A,
                                                                                                                                                                                                                                                                                                        06-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                  WO9958556-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encoding protein functionally involved in the DNA mismatch :\!\!\!\text{em} of a plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4459 TGGACTTTTTTTTTTTTTTT
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                                                                                                                                              protein associated to hereditary spastic paraplegia used is of neurodegenerative conditions.
                                                                                                                                                                                                                                                  FOND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    recombination between genomes of A.thaliana subspecies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 14 A; 4 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                      Casari G;
                                                                                                                                                                                                                                                 TELETHON.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for amplification of paraplegin
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Pred. No.
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2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hereditary spastic paraplegia;
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The present sequence is a reverse primer cmar8 used for amplification (detection of mutations in paraplegin gene exon from hereditary spastic paraplegia (HSP) patients. Detection of mutations in paraplegin gene helps in the diagnosis and treatment of various forms of HSP or other neurodegenerative conditions, such as Amyotrophic Lateral Sclerosis

Claim 4; Fig 3; 53pp; English.

diagnosis

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AAA75055
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Matches 16
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Best Local S
Matches 16
                                                                                                                                            The present PCR primer was used to amplify a human cDNA sequence, which encoded a protein with heparanase catalytic activity. The heparanase (hpa) polynucleotide is useful in gene therapy, particularly in treating tumour, inflammation or autoimmunity. Particularly, the polynucleotide is useful in modulating the bioavailability of heparin-binding growth factors, cellular responses to heparin-binding growth factors (e.g. bFGF) and cytokines (e.g. interleukin (IL)-8), cell interaction with plasma lipoproteins, cellular susceptibility to certain viral and some bacterial and protozoa infections, or disintegration of neurodegenerative plaques. The polynucleotide is also useful in wound healing (e.g. thermal, chemical or radiation burns), and in the treatment of angiogenesis, restenosis, atherosclerosis, inflammation, neurodegenerative diseases (Gerstmann-Straussler Syndrome or Creutzfeldt-Jakob disease), and some
                                                                                                                                                                                                                                                                                                                                                                 New polynucleotides encoding a polypeptide having heparanase activity, useful in wound healing and in gene therapy, particularly in treating tumor, inflammation, autoimmunity, neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                           Example
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(FRIE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       heparin-binding growth factor; cytokine; neurodegenerative plaque; wound healing; infection; burn; angiogenesis; restenosis; atherosclerosis; inflammation; neurodegenerative disease;
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                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-FEB-2000; 2000WO-US003542.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         atherosclerosis; inflammation; neurodegenerative disease;
Gerstmann-Straussler Syndrome; Creutzfeldt-Jakob disease; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; heparanase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer hpl-629 used to amplify human cDNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-SEP-2000
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4
                                                                                                                                    bacterial
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HADASIT MEDICAL RES SERVICES & DI
                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vlodavsky
                                                                                                         BP;
                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                        53; 152pp;
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                                                                                                                                    protozoa
                                                             0.2%;
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                                                                                                         C; 6
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                        7430
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Pred. No. 2.1e
0; Mismatches
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Pred. No. 2.1e+03;
0; Mismatches 2;
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                                                                 2.1e+03
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                                                                            DB 1;
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RESULT 3112
AAA63845/c
ID AAA6384
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AC AAA6384
XX

AAA63845 standard; DNA;

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AAA63845

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                                                                                                                                               AAZ65654 to AAZ69578 represent human biallelic markers from the present clinvention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOB 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                      Query Matcl
Best Local
Matches
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                                                                                                              Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                         present
                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 1735; 2745pp;
                                                                                                                                                                                                                                                                                                                                                                                                                  Novel biallelic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-1998;
23-NOV-1998;
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                                                                                  Match
                         4028 GAGAAAACAAAATGTTAT
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                                                                                                                                                                                                                                                                                                                                                                                                the human genome.
                                                                     Similarity
                                                                                                                                           invention
GAGAAAATAAACTGTTAT
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                                                                                                             BP; 9 A;
                                                      Conservative
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98US-0109732P
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18
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                                                      <u>,</u>
                                                      Score 14.8; D
Pred. No. 2.1e
0; Mismatches
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RESULT 3113
AAA63843/c
ID AAA6384
XX AAA6384
AC AAA6384
XX O4-DEC-
XX O4-DEC-
XX PCR pri
XX Human;
KW phospha
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KW phospha
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primers AAA63845-46 were used to amplify cDNA encoding full length human diacy_lglycerol kinase beta (DAGKbeta). DAGK converts diacy_lglycerol (DAG) to phosphatidic acid and attenuates DAG-dependent protein kinase C activation. Compounds that modulate the activity of DAGKbeta may be administered to a human patient for the treatment or prophylaxis of a disorder that is responsive to modulation of DAGK activity. The disorder may be a mood disorder, epilepsy, a neurodegenerative disorder, anxiety, achizophrenia, migraine, drug dependence, stroke, Alzheimer's dementia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; diacylglycerol kinase beta; DAGKbeta; diacylglycerol; DAG; phosphatidic acid; DAG-dependent protesin kinase C activation; mood disorder; epilepsy; neurodegenerative disorder; anxiety; schizophrenia; migraine; drug dependence; stroke; Alzheimer's dementia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-DEC-2000
                                                                                                   Human; diacylglycerol kinase beta; DAGKbeta; diacylglycerol; DAG; phosphatidic acid; DAG-dependent protein kinase C activation; mood disorder; epilepsy; neurodegenerative disorder; anxiety; schizophrenia; migraine; drug dependence; stroke; Alzheimer's dementia;
                                                                                                                                                                                                                      04-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human diacylglycerol kinase beta (hDAGKbeta) protein and its modulating compounds, useful for treatment of neurodegenerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-506093/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-FEB-1999;
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                      WO200047723-A2
                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                      AAA63843 standard;
                                                                                     Parkinson's disease;
                                                                                                                                                                                     PCR primer used to
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                                                                                                                                                                                                                                                                                                                                                                      19
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                                                                                                                                                                                                                                                                                                                                                                    ATTTTTCCTGGTTTGTCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 11 A; 3 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                       DNA;
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                                                                                                                                                                                     amplify cDNA ericoding full length human DAGKbeta
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                                                                                     primer;
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Pred. No. 2.
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Matches
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                                  10-SEP-1999;
26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DAG) to phosphatidic acid and attenuates DAG-dependent protein kinase C activation. Compounds that modulate the activity of DAGKbeta may be administered to a human patient for the treatment or prophylaxis of a disorder that is responsive to modulation of DAGK activity. The disorder may be a mood disorder, epilepsy, a neurodegenerative disorder, anxiety, schizophrenia, migraine, drug dependence, stroke, Alzheimer's dementia or
                                                                                                                                                                                                                                         Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New human diacylglycerol kinase beta (hDAGKbeta) protein and its modulating compounds, useful for treatment of neurodegenerative
                                                                                                                                                                                                                               pulmonary embolism;
                                                                                                                                                                                                                                                                                                                                  AAF95448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 11 A; 3 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primers AAA63843-44 were used to amplify cDNA encoding human diacylglycerol kinase beta (DAGKbeta). DAGK converts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 15; 57pp; English.
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                                                                                                       15-MAR-2001.
                                                                                                                                                                                                         Homo sapiens
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 WHITEHEAD INST BIOMEDICAL RES. MILLENNIUM PHARM INC.
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2000US-0225724P
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                                                        99US-0153357P.
                                                                                                                                                                        replace(11,G)
                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                               paternity test;
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Pred. No. 2.
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RESULT 3115
AAF96689
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Best Local
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                                                                                                                             10-SEP-1999;
26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myccardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembol pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acids comprising single nucleotide polymorphisms, us applications such as forensics, paternity testing, medicine, analysis and phenotype correlations to diseases such as diab atherosclerosis.
WPI; 2001-226749/23
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les 16; Conserv
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                                                                         WHITEHEAD INST BIOMEDICAL MILLENNIUM PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single nucleotide polymorphism #1450
                                 Gargill M,
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2000US-0220947P.
2000US-0225724P.
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                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers replace(11,T) /*tag= a
                                                                                                                                                                                                                                                                                                                       /standard_name= "single
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Nucleic acids comprising single applications such as forensics, analysis and phenotype correlati

ng single nucleotide polymorphisms, us orensics, paternity testing, medicine, correlations to diseases such as diab

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RESULT 3116
AAF96584
ID AAF9658
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Best Local
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26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                              07-SEP-2000; 2000WO-US024503
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  atherosclerosis.
                                                                     Lander ES,
                                                                                                                                                                                                                               WO200118250-A2
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                                                                                             MILLENNIUM PHARM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene
                                                                                                                                                                                                                                                                                                                                       embolism; paternity
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infarction; atherosclerosis; stroke; venous thromboembolism;
smbolism; paternity test; ds.
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2000US-0225724P.
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Pred. No. 2.
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RESULT 3117
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular disease, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPS) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                 10-SEP-1999;
26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                        Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
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                                             applications such as for analysis and phenotype
                                                                                                                                                (MILL-)
                                                                                                                                                                                                                                                                                                                                                                                               pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human gene single nucleotide polymorphism #1121.
                                    atherosclerosis.
                                                                        Nucleic
                                                                                                                                                                                                                                    07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF96360
                                                                                                                                                                                                                                                           15-MAR-2001.
                                                                                                                                                                                                                                                                                    WO200118250-A2
                                                                                                                                                                                                                                                                                                                                   Variation
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                                                                                                                                                                                                                                                                                                                                                                       sapiens.
                                                                                               2001-226749/23
                                                                                                                       ES,
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                                              acids comprising single nucleotide polymor
tions such as forensics, paternity testing,
s and phenotype correlations to diseases su
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                               WHITEHEAD I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 141;
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                                                                                                                     Gargill
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2000US-0220947P.
2000US-0225724P.
                                                                                                                                                                                                                                    2000WO-US024503
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/*tag= a
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                                                                                                                                                INST BIOMEDICAL M PHARM INC.
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                                                                                                                       Ireland
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Pred. No. 2.16
0; Mismatches
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                                                                                                                       JS,
                                                                                                                                                            RES
                                                                                                                       Bolk S,
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                                               polymorphisms, useful in
testing, medicine, genetic
eases such as diabetes and
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Example;

Page 129;

242pp;

English

The present invention provides a method of diagnosing a vascular diseas in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin genes. The sequences at a number of polymorphic sites are also provided

disease

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RESULT 3118
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Best Local :
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                                                                                                                                                                                                                             10-SEP-1999;
26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
                                                                                          Nucleic acids comprising single nucleotide polymorphisms, useful applications such as forensics, paternity testing, medicine, gene analysis and phenotype correlations to diseases such as diabetes
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                                                                                                                                                                                                                                                                                                                                                                                                                 Homo
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                                                                                  atherosclerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGCTTGAATTCATGTATGAR
                                                          Page 66;
                                                                                                                                                                  Gargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                             99US-0153357P.
2000US-0220947P.
2000US-0225724P.
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replace(11,A)
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                                                                                                                                                                                           INST BIOMEDICAL M PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotide polymorphism #241.
                                                          242pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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                                                                                                                                                                                                                                                                                                                                                      nucleotide polymorphism'
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                                                                                                                                                                    Daley
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Best Loc
Matches
PCR primers AAH28472-73 were used to amplify DNA encoding a human slit protein polypeptide, designated Zslit. Zslit is a neurite growth and development modulator, and an cellular proliferation and differentiation and immune response modulator. Zslit polypeptides and polynucleotides are useful for regenerating and directing neurite outgrowths following strokes, brain damage caused by head injuries, paralysis caused by spinal injuries, and for treating neurodegenerative diseases such as amyotrophic lateral sclerosis, Alzheimer's disease, Huntington's disease, Parkinson's disease and peripheral neuropathies, or demyelinating diseases e.g. multiple sclerosis. They are useful for lung organogenesis and repair, and thus useful for diagnosing and treating pulmonary diseases such as respiration and circulation, cystic fibrosis and asthma. They also act as a mediator of immunosuppression, and thus are useful for diagnosing and treating autoimmune diseases such as insulin dependent diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Slit protein; Zslit3; neurite growth; cellular proliferation; immune response; stroke; brain damage; paralysis; Huntington's diseas; neurodegenerative disease; amyotrophic lateral sclerosis; Alzheimer's disease; Parkinson's disease; peripheral neuropathy; demyelinating disease; multiple sclerosis; lung organogenesis; pulmonary disease; respiration; circulation; cystic fibrosis; asthma; immunosuppression; autoimmune disease; insulin dependent diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Holloway
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-DEC-1999;
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c fibrosis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00469847
                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptide, ZSLIT1, useful for treating and diagnosing insulin dependent diabetes and multiple sclerosis.
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No. 2.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   multiple sclerosis.
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Query Match Best Local S Matches 16

l Similarity 16; Conserv

Conservative

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Score 14.8; D Pred. No. 2.1e 0; Mismatches

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RESULT 3120
AAH78643
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                              PCR primers AAH78642-43 were used to amplify a gene fragment of the human mechanically sensitive potassium channel gene. The channel is activated by polyunsaturated fatty acids (particularly arachidonic acid (AA)) and by riluzole. The polypeptide is designated human TWICK-related AA-activated potassium channel (hTRAAK). The hTRAAK gene is located on chromosome 11q13. hTRAAK is involved in regulation of neuronal and muscle excitation, cardiac rhythm and secretion of hormones. Cells that express hTRAAK, designated to screen for modulators of hTRAAK activity. Such modulators are potentially useful for prevention or treatment, in humans and animals, of: cardiac and/or vascular disease; nervous system disorders associated with ischemia and anoxia; endocrinal diseases associated with anomalous hormone secretion or muscle diseases.
                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; mechanically sensitive potassium channel; riluzole; TWICK, polyunsaturated fatty acid; arachidonic acid; hTRAAK; chromosome neuronal excitation; muscle excitation; cardiac rhythm; anoxia;
                                      associated with anomalous retinal diseases. Typical
                                                                                                                                                                                                                           Disclosure;
                                                                                                                                                                                                                                                    cardiovascular diseases and polyunsaturated fatty acids.
                                                                                                                                                                                                                                                                              New mechanically sensitive
                                                                                                                                                                                                                                                                                                                                 Lazdunski M,
                                                                                                                                                                                                                                                                                                                                                                                                                14-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                           20-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200168670-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hormone secretion; cardiac disease; vascular disease; ischem
nervous system disorder; endocrinal disease; muscle disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rheumatoid arthritis
                          neurodegeneration
                                                                                                                                                                                                                                                                                                                                                                                     14-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    retinal disease; epilepsy; cardiac arrhythmia; neurodegeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH78643;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH78643 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2530 ACAGCAGATGAGCTCCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for mechanically sensitive potassium channel gene
                                                                                                                                                                                                                         Page 15; 37pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                              CENT
                                                                                                                                                                                                                                                                                                                                                                                     2000FR-00003264.
                                                                                                                                                                                                                                                                                                                                                                                                                2001WO-FR000758
                                                                                                                                                                                                                                                                                                                                 Lesage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                         examples are
                                                                                                                                                                                                                                                               potassium channel, useful for
in drug screening, is activat
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Pred. No. 2
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                                       epilepsy,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2
                                         cardiac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21;
                                         arrhythmia
                                                                                                                                                                                                                                                                              treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TWICK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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XX (CRS)
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                                                 Ś
                                                                                                             Matches
                                                                                                                                    Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                              PCR primers AAH78639-40 were used to amplify a gene fragment of the human mechanically sensitive potassium channel gene. The channel is activated by polyunsaturated fatty acids (particularly arachidonic acid (AA)) and by riluzole. The polypeptide is designated human TWICK-related AA-activated potassium channel (hTRAAK). The hTRAAK gene is located on chromosome 11g13. hTRAAK is involved in regulation of neuronal and muscle excitation, cardiac rhythm and secretion of hormones. Cells that express hTRAAK, designated to screen for modulators of hTRAAK activity. Such modulators are potentially useful for prevention or treatment, in humans and animals, of: cardiac and only vascular disease, nervous system
                                                                                                                                                                                                                                                                                                      disorders associated with ischemia and anoxia; endocrinal diseases associated with anomalous hormone secretion or muscle diseases; and retinal diseases. Typical examples are epilepsy, cardiac arrhythmis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polyunsaturated fatty acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New mechanically sensitive potassium channel, useful for treating cardiovascular diseases and in drug screening, is activated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-590037/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lazdunski M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-MAR-2001; 2001WO-FR000758.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hormone secretion; cardiac disease; vascular disease; ischemia; nervous system disorder; endocrinal disease; muscle disease; retinal disease; epilepsy; cardiac arrhythmia; neurodegeneration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; mechanically sensitive potassium channel; riluzole; TWICK; polyunsaturated fatty acid; arachidonic acid; hTRAAK; chromosome 11q13; neuronal excitation; muscle excitation; cardiac rhythm; anoxia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 15; 37pp; French.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer for mechanically sensitive potassium channel
                                                                                                                                                                                                                                                                            neurodegeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CNRS ) CNRS
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                                                                                                             16;
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                                                                                                                                       Similarity
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                                              GCCCGGCCCGGGATCCCG
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                                                                                                                                                                                                                     BP; 3
                                                                                                             Conservative
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                                                                                                                                    0.2%;
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                                                                                                                                                                                                                        C; 9
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                                                                                                                                       Score 14.8;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                               arrhythmia
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RESULT 3122

Arachidonate 12-lipoxygenase polymorphism containing

DNA fragment

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                                                                                                   RESULT 3123
AAH62539
                                                                                                                                                                                                                      Query Match
Best Local 9
                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                               DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Single nucleotide polymorphism; SNP; human; heart disease; paternity testing; forensic (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-MAY-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH62114 standard; DNA; 21
                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 29; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-367705/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-NOV-1999;
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                                    12-SEP-2001
                                                                                        AAH62539 standard; DNA; 21
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                                                                                                                                                                                                           Conservative
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                                     (first entry)
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                                                                                                                                                                                                                                                           7 A;
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replace(11, A)
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                                                                                                                                                                                                                                                             C; 3
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                                                                                          ВP
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                                                                                                                                                                                                                Score 14.8; Db 1,
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0; Mismatches
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U;
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ARESULT 3124
AAF64065
ID AAF6406
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XX AAF6406
XX O6-APR-
XX Human;
XX Human;
XX Homo sa
XX HOZOO10
XX J0-JUN-
XX 30-JUN-
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                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                        DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
                                                                                                                                                                                                                                                                                                                                                                                      Sequence
 30-JUN-2000; 2000WO-US018308.
                                                                                                    Human; lipoprotein
                                                                                                                                                           06-APR-2001
                                                                                                                                                                                     AAF64065
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphism; SNP; human; cancer; heart disease; paternity testing; forensic science;
                                                    WO200102606-A2
                                                                             Homo sapiens
                                                                                                                                                                                                              AAF64065 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 64; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-NOV-1999;
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                                                                                                                                                                                                                                                                                                         6216 AAAAGGTGGGAAAGGAGA
                                                                                                                                                                                                                                                                                4
                                                                                                                                                                                                                                                                                                                                   l Similarity
16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                      BP; 10
                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                           (first
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                                                                                                     lipase;
                                                                                                                                                                                                                                                                                                                                                                                      A; 1 C; 8
                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                             0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BIOMEDICAL
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                                                                                                    LPL; stenosis;
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Pred. No. 2.1e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                     G; 2
                                                                                                                                                                                                                                                                                                                                                                                     T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                              2.1e+03
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AAH78090/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to detecting a genetic predisposition in a human subject for non-responsiveness to statin drug treatment, involving amplifying nucleic acids including a non-coding or untranslated region within the 3' end of the human lipoprotein lipase (LPL) gene from a tissue sample. The method is useful for determining which partients suffering from coronary artery dypass graft (CABG) patients, will likely not respond positively to statin drug treatment with respect to stenosis of a coronary artery or bypass graft
                                                                                                                                                                                                                                                                                         Breast amplified G protein coupled receptor; BCA-GPCR; chromosome 1q44; BCA-GPCR-1; BCA-GPCR-2; BCA-GPCR-3; BC signal transduction; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genetic testing for determining non-responsiveness to statin drug in patients of a coronary artery disease, involves analyzing amplification products for homozygosity for a variant allele in the human lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ,02-JUL-1999;
                                             WPI; 2001-570865/64.
                                                                                                         (POWE/)
                                                                                                                                                      14-MAR-2000;
11-APR-2000;
                                                                                                                                                                                           13-MAR-2001; 2001WO-US008020
                                                                                                                                                                                                                                          WO200168704-A2
                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                         Primer for breast amplified G
                                                                                                                                                                                                                                                                                                                                                                  26-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                          AAH78090;
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                                                                     Powers
                                                                                                                                                                                                                   20-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                  AAH78090 standard;
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                                                                                                         YANG J.
                                                                                                                               TULARIK INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
Similarity 88.9%;
                                                                                            CUTLER G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCACTGTTTCTTGTAAGT
                                                                   Yang J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                      2000US-00524730
2000US-00546986
                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                     Cutler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 C; 5 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                    21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1;
Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                          protein coupled receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yang
                                                                                                                                                                                                                                                                                                     BCA-GPCR-3; BCA-GPCR-4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                  breast cancer;
                                                                                                                                                                                                                                                                                                                                           (BCA-GPCR) -2.
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Four nucleic acids encoding breast amplified G protein coupled receptors (BCA-GPCRs), useful for identifying modulators of G-protein coupled receptor signal transduction which can be used in the treatment of cancer

the treatment of cancer

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RESULT 3126
AAH89072
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primers AAH78089-90 were used to amplify cDNA or DNA encoding breast amplified G protein coupled receptor (BCA-GPCR)-2. BCA-GPCRs are amplified and/or overexpressed in breast cancer cells. The BCA-GPCRs are located at chromosome 1q44, in the following orientation (starting from the centromere end): BCA-GPCR-1 (3'-5' orientation), BCA-GPCR-2 (5'-3' orientation), BCA-GPCR-3 (3'-5' orientation), BCA-GPCR-3 (3'-5' orientation), and BCA-GPCR-4 (5'-3' orientation), BCA-GPCR-3 (3'-5' orientation), and BCA-GPCR-4 (5'-3' orientatifying modulators of G-protein coupled receptor signal transduction. The modulators and antibodies against the G protein coupled receptors are useful for pharmacological modulation of signalling pathways, e.g. in cancer cells such as breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             such as breast cancer
The present invention relates to human oligonucleotides comprising single nucleotide polymorphic site (SN: AAH88797-AAH89219). The presented is one such oligonucleotide. The oligonucleotides can be usequence is one such oligonucleotide. The oligonucleotide with be used to the companion of polymorphisms with
                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                        plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                    paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphic; SNP; forensic science;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human polymorphic oligonucleotide L21952 fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH89072 standard;
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                                                                                         New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly and also in forensics and paternity testing.
                                                                                                                                                                                                                                                                                           17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                        Variation
                                                                                                                                                                         Au K,
                                                                                                                                                                                                   (GLAX )
                                                                                                                                                                                                                                                                                                                     WO200134840-A2
                                                                  Claim
                                                                                                                                                                                                                                          10-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3894
                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
                                                                                                                                               2001-335945/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7;
                                                                                                                                                                                                                                                                                                                                                                                                                                        breeding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           l Similarity
16; Conserv
                                                                                                                                                                      Chen J,
                                                                                                                                                                                                  GLAXO GROUP LTD.
AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTGGAGTTACTTTCATAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                         eding; ds.
                                                                Page 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54; 68pp;
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                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers replace(11,a) /*tag= a
                                                                                                                                                                         Patil N,
                                                                                                                                                                                                                                                                                                                                           /standard_name= "single nucleotide polymorphism"
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                                                                English
                                                                                                                                                                         Thomas
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Pred. No. 2
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U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     genetic mapping; animal breeding;
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             ne present
be used in
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RESULT 3127
AAF89455
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                          The present invention describes a method of inentifying the presence of a cc plurality of genetic markers in a subject, involving generating extension cc products using PCR primers flanking the plurality of markers, separating ct the extension products depending on their melting temperatures, and can lysing them to determine the presence or absence of each genetic marker. This can be used in the diagnosis of genetic diseases, including cmarker. This can be used in the diagnosis of genetic diseases, including csickle cell disease, phenylketonuria, galactosaemia, fragile X syndrome, csickle cell disease, phenylketonuria, galactosaemia, fragile X syndrome, csickle cell disease, cystinuria, methylmolonic acidaemia, urea cycle maturity onset diabetes, cystinuria, methylmolonic acidaemia, urea cycle disorders, hereditary fructose intolerance, hereditary haemochromatosis, coliperase, acaptonuria, hypolactasia, Baker's disease, tyrosinaemia, Wilson's cdisease, acaptonuria, hypolactasia, Baker's disease, argininaemia, cc ancitryppsin deficiency, Duchenne muscular dystrophy, Marfan's syndrome, neurofibromatosis, osteogenesis imperfecta, retinoblastoma, Freidrich's cataxia, haemoglobinopathies, Leber's hereditary optic neuropathy, MCAD, canxiaria, haemoglobinopathies, Leber's hereditary optic neuropathy, MCAD, canxiaria, cantitrypsin deficiency cetinitis pigmentosa, Bloom syndrome, Franconi anaemia
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting multiple genetic markers in one assay, useful to simultaneously detect a number of genetic disorders, comprises generating extension products and separating them on the basis of melting behavior is.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genetic marker; genetic disease diagnosis; cystic fibrosis; haemophilia; sickle cell disease; muscular dystrophy; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          phenotypic traits, assisted breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genetic marker PCR primer SEQ ID NO: 44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF89455 standard;
            Canavan's disease, retinitis pigmentosa, Bloom or Neimann Pick disease. The present sequence i of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-329096/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunlop CLM, Weisel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  retinoblastoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                       Claim 44; Page 36; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , genetic mapping of phenotypic traits and marker of animals and crop plants
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                                                                                                                                                                                                                                                                                                                                                          New nucleic acids with endonuclease activity, such as ribozymes and nucleozymes, for modulating gene expression in a plant, mammalian, bacterial or fungal cell.
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(UYYA )
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31-MAR-2000;
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AAS12385 standard;
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                                                                                 The invention relates to nucleic acid molecules with endonuclease activity, which are particularly useful for cleavage of RNA or DNA. The nucleic acids are used in a pharmaceutical composition and are used to modulate expression of a gene in a plant, mammalian, bacterial or fungal cell. They are used to cleave a separate nucleic acid, preferably RNA. The nucleic acids are used to inhibit gene expression and/or cell proliferation, and can be used to treat a disease or condition. More than one nucleic acid can be independently targeted to the same or different sites in a cell. The nucleic acids may be used to study DNA. The modifications to the nucleic acids optimises their catalytic activity and can maintain or enhance their activity. They exhibit a high degree of specificity for RNA. The present sequence represents the Class I-XII ribozyme substrate, used in an example which demonstrates the method of
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4911 TGGAGAAAGCATCAGGAC 4928

RESULT 3131

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RESULT 3130
ABA10117
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                              S
                                                                                                                                                              The present invention relates to a method for sorting genes. The method comprises producing first double stranded (ds) cDNA from mRNA by reverse cranscription using a poly-T primer. The ds cDNA is then digested with a creativation enzyme that generates cohesive ends with overhanging single creativation enzyme that generates cohesive ends with overhanging single creativation products are ligated to a set of ds DNA oligonucleotide, and the city of the daptor has at one end, a sequence complementary to a gossible overhang and the other end a primer-template sequence specific creative for the adaptor complementary sequence, and between these two ends the creative for the adaptor complementary sequence. An eligated cDNA moleculas are amplified in separate PCR assays, using for each a primer that anneals to comply and a second primer, from a set that anneals to the CDNA specific primer-template sequences. Amplicons are finally sorted into non-credundant groups defined by the specific primer that annealed to the primer-template sequences and thus primed PCR. The method is useful for primer-template sequences and thus primed PCR. The method is useful for compressed-gene transcript in the original sample is represented by the swn subgroup. The method is also useful for isolation, identification or analysis of genes, analysis and diagnosis of diseases, for studying cell differentiation and in gene therapy. The present sequence was used to illustrate the method of the present invention
                                                                   Matches
                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sorting genes into non-redundant groups, useful ediagnosis and in gene therapy, by amplifying cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gene sorting; PCR primer; disease diagnosis; disease analysis; cell differentiation; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       selective adaptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-626451/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tail primer #110 from primer set 256 used in gene sorting method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABA10117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABA10117 standard;
                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Fig 13; 67pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ulanovsky L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-MAR-2001; 2001WO-US009392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200175180-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (QBIQ-) QBI ENTERPRISES LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-OCT-2001
                                 6888 GTTGGTGCTCTCCCTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ||||| || ||||||||||| 18 TGGAGTAACCATCAGGAC 1
                                                                 Similarity 88.1
                                                                                                                                     21
                                                                                                                                   B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-00538709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mugasimangalam R,
                                                                                                                                     N
                                                                                                                                     ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                 0.2%;
                                                                                                                                       œ
                                                                                                                                     C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                 6905
                                                                   <u>,</u>
 18
                                                                                                                                       G; 7 T; 0 U;
                                                                                   Score 14.8; DB 1
Pred. No. 2.1e+03
                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Einat P,
                                                                                                                                       0 Other;
                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zezin-Sonkin D,
                                                                                                 Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .g. for gene isolation, fragments attached to
                                                                     Indels
                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shlomit
                                                                     Gaps
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AAD29834/c

ID AAD29834 standard; D)

XX AC AAD29834;

XX AC AAD29834;

XX 17-MAY-2002 (first |
XX X 2002983 |
XX X 2002983 |
XX X X 2002983 |
XX X 20020198509-A2.

XX W0200198509-A2.

XX W0200198509-A2.

XX 21-JUN-2001; 2001W0-XX 21-JUN-2000; 2000US-YA PAF

XX 21-JUN-2001; 2001WO-PAF

XX 21-JUN-2001; 2000US-PAF

XX 21-JUN-2001; 2000US-PAF

XX 
XXXXXXXXXXXXXXX
                                                                                                                                                                                                                                                                                                                                  RESULT 3132
ABL49744
                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a transgenic plant comprising heterologous DNA coding for eukaryotic thioredoxin reductase integrated into its nuclear or plastid genome and use of thioredoxin reductase for separating the starch and protein components of grain in a milling process. Transgenic plant is used for separating the starch and protein components of grain in a milling process. Transgenic plant may be used to produce thioredoxin reductase at elevated levels. Delivery of thioredoxin reductase eliminates the need to develop exogenous sources for addition during processing. Secondly, physical disruption of seed integrity is not necessary to bring the enzyme in contact with the storage or matrix proteins of the seed prior to processing or as an extra processing step.

The present sequence is Arabidopsis NADPH dependent trx reductase gene
                                                 Human;
plant;
Oryza sativa
                                                                                                                                                                                            29-MAY-2002
                                                                                                                                                                                                                                                   ABL49744;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 4; Page 52; 86pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Transgenic plant coding for eukaryotic thioredoxin reductase at elevated levels useful for separating the starch and protein components of grain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transgenic plant; thioredoxin reductase;
                                                                                                                                        Rice PCR
                                                                                                                                                                                                                                                                                                       ABL49744 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lanahan MB, Desai NM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUN-2000; 2000US-00598747.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUN-2001; 2001WO-EP006918.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        milling process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplifying PCR
                                                 grass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16;
                                                 interferon alpha; interferon omega; prolamin; fusion protein; grass plant; tea; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             process; enzyme; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CATCCCTAAGATCGACGT 368
                                                                                                                                  primer SEQ ID NO:6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                CAACCCGAAGATCGACGT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ۸
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dependent trx reductase gene amplifying primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer
                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gasdaska PY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G; 6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 1;
Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    starch; protein; grain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRF2B
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ABK70370
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a true grass plant in which a base sequence encoding a human interferon protein, with a base sequence encoding a signal amino acid sequence causing accumulation of the stored protein of albumen to the protein body connected upstream, is recombined expressably to a genomic DNA. The grass plant can be used for the creation of a useful application of tea. The present sequence represents a PCR primer which is used in an example from the present invention
                                                                                                                                                                                                                                                  insulin-like growth factor binding protein-2; hormone-regulated tumour; breast cancer; prostate cancer; IGF-1-sensitive cancer; apoptosis; hormone-responsive cancer; hormonal withdrawal; oligodeoxynucleotide; ODN; endocrine tumour therapy; ss.
                                                                                       Gleave
                                                                                                                                   14-SEP-2000; 2000US-0232641P
                                                                                                                                                          13-SEP-2001; 2001WO-US028748
                                                                                                                                                                                  21-MAR-2002.
                                                                                                                                                                                                       WO200222642-A1.
                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                               Hormone-regulated cancer; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                    Synthetic antisense IGFBP-2-oligodeoxynucleotide (ODN)
                                                                                                                                                                                                                                                                                                                                             15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                         ABK70370 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 4; 22pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUL-2000; 2000JP-00207230.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUL-2000; 2000JP-00207230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JP2002017187-A.
                                                                 WPI; 2002-339861/37
                                                                                                                                                                                                                                                                                                                                                                    ABK70370;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-263239/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NICH-) JAPAN CHEM RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                              6317 GGCTACTGTTGCTGGGAA 6334
                                                                                       Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
                                                                                                              VIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                GGCTAATGTTGGTGGGAA 21
                                                                                      Satoshi K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              incorporated human interferon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                              BRITISH
                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 2 C; 8 G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.2%;
                                                                                                              COLUMBIA.
                                                                                       Nelson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CO LID
                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>..</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                       Rennie
                                                                                       PS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                               IGFBP-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Example 1; Page 13; 36pp; English

Composition for treating hormone-regulated cancer, particularly of prostate or breast, comprises oligonucleotide antisense to insulin-like growth factor binding protein-2.

문 5

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ABS52265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a new composition for treating hormone regulated cancer. The composition comprises an antisense oligonucleotide that inhibits expression of IGFBP-2 (insulin-like growth factor binding protein-2). The molecules of the invention are used to delay progression of hormone-regulated tumours, particularly of breast or prostate, to the hormone independent state, to delay metastatic progression to the bone of IGF-1-sensitive cancers and to treat hormone-responsive cancers by inducing apoptosis, after hormonal withdrawal. The present nucleic acid sequence represents one of a collection (ABK70313-ABK70375) of antisense IGFBP-2-oligodeoxynucleotides (ODN) that were used in the invention for
                                           The invention relates to producing multicellular plants, organs or tissues transformed on their plastome, comprising altering/disrupting the function of a gene in a plastid genome for producing a selectable phenotype and selecting plants with plastids expressing the phenotype, transforming the plastid genomes of selected plants with a transformation vector with a restoring sequence for restoring function and separating transformed plants. This method is useful for producing multicellular plants, organs or tissues transformed on their plastome and for selection of antibiotics and herbicide resistance genes. Sequences ABS52237-ABS52269 represent PCR primers used to amplify plant vector genes of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Plant; PCR;
herbicide re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS52265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IGFBP-2-oligodeoxynucleotides (ODN) that we prostate and other endocrine tumour therapy
                                                                                                                                                                                                                                                                       Transforming multicellular plants, by altering the function of a plastid gene, selecting plants expressing altered phenotype, transforming plants with a vector capable of restoring function and separating transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JAN-2001; 2001DE-01002389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200257466-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plant vector
                                                                                                                                                                                                                            Example 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-JAN-2002; 2002WO-EP000481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCACTAGCAGCAGCAGCA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCAGCAGCAGCAGCA
                                                                                                                                                                                                                                                                                                                                                                   Huang F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                resistance
                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; ss; plastome; plastid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR
                                                                                                                                                                                                                                                                                                                                                                                                      GENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                            38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer #29.
                                                                                                                                                                                                                                                                                                                                                                      Klaus S,
                                                                                                                                                                                                                        56pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene.
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                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
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Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                      Muehlbauer S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genome; antibiotic;
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                                                                                                                                                                                                                                                                                                                                                                      Herz S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2
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RESULT 3135
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Best Local S
Matches 16
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                                                                                                                                                                                                                                            26-OCT-2000

26-OCT-2000

26-OCT-2000

27-OCT-2000

27-OCT-2000

31-OCT-2000

01-NOV-2000

02-NOV-2000

02-NOV-2000

02-NOV-2000

19-UN-2001
                                                                                                              Gangolli EA,
Vernet CAM,
Malyankar U,
          Isolated NOVX polypeptides and nucleic acid molecules useful itreating, preventing, diagnosing and researching pathological in humans with a NOVX-associated disorders, e.g. cancer, stroiplisheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C5-NOV-2002
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                                                                                                                                                                                                                     15-FEB-2001;
02-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                 26-OCT-2001; 2001WO-US050925
                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; NOVX; pathological condition; NOVX-associated disorder;
                                                                                                                                                                   (CURA-)
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                                                                           2002-590673/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
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2000US-0243641P.
2000US-0243681P.
2000US-0243681P.
2000US-0244943P.
2000US-0244995P.
2000US-0244995P.
2000US-0245315P.
2000US-0245315P.
2000US-0245316P.
2000US-026294P.
2000US-026294P.
2001US-026294P.
2001US-026294P.
2001US-02685P.
2001US-0276565P.
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                                                                                                               Spytek KA,
Shenoy S, N
Stone D, N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                    Anderson
                                                                                                                                                                     CORP
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                                                                                                               ,, Gilbert J, Casman S, Blalock A, Li L;
Mishra V, Furtak K, Gerlach V, Edinger
Millet I, Smithson G, Gunther E, Padiga
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Pred. No. 2
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                                                                                                                  dinger S;
Padigaru
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                                                    for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        psoriasis;
                                      conditions
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Example 3;

Page

170;

236pp; English.

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with

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RESULT 3136
ABK9120
ID ABK9120
XX
AC ABK9120
XX
DE Human
XX
Human;
KW Human;
KW Jaran;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          syndrome, cirrhosis, transplantation disorders, pancreatitis, obesity, diabetes, autoimmune disease, renal artery stenosis, interstitial nephritis, glomerulonephritis, polycystic kidney disease, cataract, Alzheimer's disease, acoustic trauma, cancer, infertility, cardiomyopathies, atherosclerosis, hypertension, congenital heart defects, scleroderma, endometriosis, haemophilia, dementia, stroke, Parkinson's disease, Huntington's disease, epilepsy, multiple sclerosis, anxiety, pain, leukaemias, hypothyroidism, psoriasis, acne, wounds and asthma. They are also useful for the manufacture of a medicament for treating a syndrome associated with a human disease, specifically a Novx-associated disorder. They may also be useful in therapeutic applications including protein therapy, as small molecule drug targets, as antibody targets, as diagnostic and/or prognostic markers, in gene therapy, as research tools and in tissue research tools and in tissue research.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; 88; Ulip; CRMP; collapsin response mediator protein; PCR; Unc-33-like protein; neurodegenerative disease; Alzheimer's disease; paraneoplastic neurodegenerative disease; PND; myelination; demyelination; remyelination; myelin disorder; multiple sclerosis; autoimmune neurodegenerative disorder; HTLV-1 associated myelopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a new polypeptide that comprises any of 17 fully defined sequences of 43-990 amino acids given in the specification. The NOVX polypeptide, nucleic acid and antibody of the invention are useful for treating or preventing a pathological condition in humans with a NOVX-associated disorder, e.g. Von Hippel-Lindau
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human T lymphocyte virus 1; reverse proteolipid protein; PLP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      research tools and in tissue regeneration. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention to amplify one of the 17 novel proteins of the invention
            WPI; 2002-627172/67
                                                                                                                                                                                                                                                                                                                                           09-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                 09-NOV-2001; 2001US-00986632.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2002119944-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human proteolipid, PLP, protein RT-PCR primer #1.
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                                                                 Aguera M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-AUG-2002
                                                                                                                                                                                                                                                                                (AGUE/)
                                                                                                                                                                                                                    (CHAR/)
                                                                                                                                                                                                                                                    (BELI/)
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                                                                                                                      BELIN M.
CHARRIER B.
HONORAT J.
RICARD D.
ROGEMOND V.
                                                                                                                                                                                                                                                                                AGUERA M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 BP; 8 A;
                                                              Belin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                           2000US-0246751P
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                                                                 Charrier E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
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                                                                 Honorat J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcriptase PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.1e+03
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                                                                 Ricard
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                                                              Rogemond V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
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PIXBXRRXRXX
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ABL57072/c
ID ABL57072 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC myelin disorders, comprises administering to a patient an effective (collapsin response mediator protein) protein, a nucleic acid coding for CC (collapsin response mediator protein) protein, a nucleic acid coding for CC (lip)(CRMP, an antisense sequence capable of specifically hybridising with CC the nucleic acid, an antibody directed against Ulip/CRMP, or an aptamer CC capable of binding ULIP/CRMP, and a pharmacologically acceptable carrier. CC Also included are methods of diagnosing a myelin disorder in a subject, cidentifying agents useful for the prevention or treatment of myelin CC disorders, using the Ulip/CRMP proteins/nucleic acids, agents capable of CC modulating the function or expression of the proteins (increasing or CC decreasing), and a method for identifying an endogenous agent as a CC therapeutic target for the prevention or the treatment of myelin CC disorders. The agents are useful for preventing or treating a myelin CC disorder such as multiple sclerosis or HTIV-1 (human T lymphocyte virus C lassociated myelopathy and neurodegenerative diseases, Alzheimer's C disease, paraneoplastic neurodegenerative diseases (PND), autoimmune C theorogenes of myelination. Asswering are involved in articles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prevention or treatment of myelin disorders, such as multiple sclerosis, by administering an agent selected from a Ulip/CRMP protein, a nucleic acid coding for the protein, or an antibody directed against protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      theprocesses of myelination, demyelination and remyelination. Antibodies to a Ulip/CRMP protein are useful for diagnosting a myelin disorder. The present sequence is a reverse transcriptase (RT)-PCR primer for proteolipid protein (PLP, an oligodendrocytic marker protein) used as a control in an experiment to detect mRNA encoding Ulip proteins
                             (UYRQ ) UNIV
                                                             29-AUG-2000;
30-MAR-2001;
                                                                                                                                               07-MAR-2002.
                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                   Molecular beacon;
                                                                                                                                                                                                                                                                                                                                                                   Molecular beacon target sequence (single mismatch).
                                                                                                                                                                                                                                                                                                                                                                                                      22-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Page 8;
                                                                                                              29-AUG-2001; 2001WO-US041941
                                                                                                                                                                                WO200218951-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL57072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention relates to a new method in disorders, comprises administer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 5 A; 3 C; 8 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                             2000US-0228728P
2001US-0280350P
                             ROCKEFELLER
                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                /*tag= a
/note= "mismatch site'
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                                                                                                                                                                                                                                                                                                                                   fluorophore;
                                                                                                                                                                                                                                                                                                                                                                                                      entry)
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                                                                                                                                                                                                                                                                                                                                   nanoparticle; nucleic
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                                                                                                                                                                                                                                                                                                                                   acid
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                                                                                                                                                                                                                                                                                                                                   detection; ss
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Dubertret B,

Calame M, Libchaber A;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC molecular beacon comprising an oligonucleotide probe (see ABL57069)
CC covalently attached at the 3' end to fluorescent dye and at the 5' end to
CC a nanoparticle. In the native state, the probe forms a hairpin
CC conformation with hybridised termini. The proximity of the fluorophore
CC and quencher (gold nanoparticle) in the molecular beacon results in
CC little or no detectable fluorescence. Upon hybridisation of the central
CC complementary stretch of the probe to a target sequence, such as the
CC present sequence, the hairpin undergoes a conformational change resulting
CC in an increase in fluorescence, the extent of which is proportional to
CC the amount of target sequence present. Experiments with the present
CC sequence and a perfectly-matched target (see ABL57071) showed that
CC relates generally to the use of metal surface quenchers such as particles
CC relates for high sensitivity applications in, for example, detection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-404569/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide array; hydrophobic attachment layer; diagnosis; cholesterol; genetic analysis; DNA chip; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD22716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD22716 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21
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   Attaching oligonucleotide on support by fluidizing hydrophobic attachment
                                                                                                                                                                                                                                         14-APR-2000; 2000KR-00019557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fluorescent-oligonucleotide Flu. 4 used for hybridisation.
                                                             WPI; 2002-026037/03
                                                                                                                                                                                                                                                                                                   29-MAR-2001; 2001WO-KR000514.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 14 A; 4 C; 2 G; 1 T; 0 U; 0 Other;
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FFX8X333333333333333X8 layer applied on support, spotting oligonuclectide solution having one of its ends bonded to hydrophobic group on layer and solidifying layer.

Example 1; Page 12; 32pp; English

The invention relates to a method for attaching oligonucleotide to a solid support and an oligonucleotide array. The oligonucleotide array is prepared by applying hydrophobic attachment layer onto a support which can be solidified under certain conditions, spotting certain portions of said attachment layer with aqueous oligonucleotide solution in which hydrophobic groups are bonded to 3' or 5' terminals and solidifying said attachment layer. The method is useful for attaching a oligonucleotide (preferably a PCR amplified product from a PCR with its 5' terminal bonded to a hydrophobic group) onto support. The method is useful in the area of genetic diagnosis and analysis, and DNA chips which are based on hybridisation. The present sequence is fluorescent oligonucleotide useful in hybridisation. hybridisation. The in hybridisation

Sequence 21 BP; 11 A; 1 C; 7 G; 2 T; 0 U; 0 Other;

S 밁 Matches Query Match Best Local (2904 TGCCTTGTTTCCTTCTAT 2921 16; 20 Similarity TGACTTCTTTCCTTCTAT Conservative 0.2%; ω 0; Mismatches Score 14.8 Pred. No. 3 14.8; L. No. 2.1e+03; DB 1; 2 Length Indels 21 0, Gaps 0

RESULT 3139 ABS98296/c ABS98296 standard; DNA; 21 BP 23-DEC-2002 (first entry) (LTF) gene polymorphic sequence

KW cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
KW cytochrome P450 A2; CYP4501A2; Cytochrome P450 02E; CYP45002E1; LTF;
KW cytochrome P450 A2; CYP4501A2; Cytochrome P450 02E; CYP45002E1; LTF;
KW cytochrome P450 A2; CYP4501A2; Cytochrome P450 02E; CYP45002E1; LTF;
KW cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP450 AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW glutathione-S-transferase 12; GST12; histamine-N-methyl transferase; NMT;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NMT;
KW MDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7;
KW UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; UPA;
KW UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; UPA;
KW multidrug resistance associated protein 3; cancer; prosetate;
KW multidrug resistance associated protein 3; cancer; prosetate;
KW multidrug resistance associated protein 3; cancer; prosetate;
KW acettral nervous svatem; pulmonarv- immunological xND: single central nervous system; pulmonary; immunological; SNP nucleotide polymorphism.

WO200257410-A2

25-JUL-2002.

28-NOV-2001; 2001WO-US044838

28-NOV-2000; 2000US-00724389

DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC expression, mutation or underexpression, which may be used in diagnosing CC and/or treating the disorders. The nucleic acid molecules comprising the CC polymorphic sequences contained in CYP4501A1, CYP94501A2, CYP4502E1, CC ARNY, EPHX2, GST12, NNWY, NQO2, NR112, STM, UGT2B3, UGT2B3, UGT2B15, AHR, CMDR1 and/or MDR3 are useful for screening individuals for altered drug CC metabolism. The polymorphic sequences contained in CYP4501A1, CYP94501A2, CRHR, MDR1 and/or MDR3 may also be used to screen individuals for CC susceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are CC used to screen for altered cardiovascular function, in CA2 for altered contral convolus system function, in FLAP and HNMT for altered derine CC immunological or haematological function, in LTP for altered immunological or haematological function, in CHMR3, CHMR4 or CHMR5 for altered central and CC immunological function, in CHMR3, CHMR4 or CHMR5 for altered central and CC peripheral nervous system function. The present sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC cytochrome P450 0221 (CYP4500221), adrenergic receptor betal (ADBRI), CC cytochrome P450 0221 (CYP4500221), adrenergic receptor betal (ADBRI), CC (ARWT), cathepsin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding CC inhibitor (DBI), epoxide hydroxylase 2 (EPH22), 5-lipoxygenase activating CC protein (FLAP), glutathione-S-transferase 12 (GST12), histamine-N-methyl CC transferase (HNWT), (kallikrein 2) KLKz, nicotinamide -N-methyl CC transferase (NNWT), NADPH quinone oxidoreductase 2 (NQO2), cultotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B4 (CYP3B15), urokinase receptor (uPA), multidrug resistance 1 (CMRR), clarase receptor (NRII2), or acetylcholine muscarinic CC receptor 1, 2, 3, 4, or 5 (CHMR), CHMR3, CHMR4 or CTHR5) sequence. CC The polymorphisms in the human genes cited in the invention are useful as CC genetic linkage markers for locating and characterising the genes that CC are responsible for a specific traits within the genome and eventually CC identifying the genes responsible for a variety of disorder-related CC expression multiation or undergraphs with man be used in diagnosing CC expression or undergraphs or wariety of disorder-related CC expression multiation or undergraphs which may be used in diagnosing CC expression or undergraphs of the mass of the diagnosing constitutive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                  Bacteriophage lambda; loop mediated isothermal amplification; LAMP; polynucleotide synthesis; single nucleotide polymorphism; SNP; identification; primer; ss.
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                     Bacteriophage lambda Synthetic.
                                                                                                                                                                                                                             27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                Bacteriophage
                                                                                                                                                                                                                                                                                                                      ABL51321
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                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA
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                                                                                                                                                                                lambda related LAMP primer SEQ ID NO:55
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Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymucleotide having a loop at one end and containing two copies of the template; (b) using a second (loop) primer set to initiate complementary chain synthesis at a different position; (c) repeating the process; and (d) carrying out DNA polymerase catalysed complementary chain synthesis. The method is simple and efficient for polymucleotide amplification, allowing easy identification of the presence of single nucleotide polymorphisms (SNP). The method is isothermal, produces a high degree of amplification in a short time, and since it uses primers hybridising to several different sequences on the template it is highly accurate, without the amplification of irrelevant sequences which is a problem in conventional methods. The present sequences which is a problem in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for polynucleotide synthesis by loop-mediated isothermal amplification, (LAMP). The method comprises: (a) identifying a target sequence on the template and a set of primers (inner and outer primer) hybridised to it and annealed to form a single-stranded polynucleotide with a loop at each end, followed by self-primed strand displacement complementary chain synthesis to form a two-stranded polynucleotide with a loop at each end, followed by self-primed strand displacement complementary chain synthesis to form a two-stranded polynucleotide having a loop of the lamburge of the l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28 MAR - 2002
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rapid isothermal polynucleotide synthesis using loop-mediated amplification for simple detection of single nucleotide polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200224902-A1
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                                                                                                                                                                                                   4368 ACAGGCTGGGGAATTTTG 4385
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Pred. No. 2
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X P X O X X W W W W W W W W X D X D X A X X
                                                                           NOV15 forward PCR primer #2.
                                                                                      21-OCT-2002
                                                                                                          ABQ94035 standard; DNA; 21 BP
                                                                                     (first entry)
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Human; NOV; cytostatic; Cardiant; Antiinflammatory; Immunosuppressive; Antiallergic; Haemostatic; Anti-HIV; Antidiabetic; Anorectic; Antiasthmatic; Nephrotropic; Hapatotropic; Neuroprotective; Nootropic; Antibacterial; Virucide; Antiparasitic; Relaxant; Anticonvulsant; Gene Therapy; NOV; cancer; heart disease; inflammation; autoimmune disorder; allergy; blod disease; Inflammation; asthma; IgA nephropathy; cirrhosis; arthritis; Alzheimer's disease; infection; stroke; muscular dystrophy; epilepsy; wasting disorder; PCR sapiens.

WO200255704-A2

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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Padigaru M, Li L, Zerhusen Zhong M, Gangolli EA, Burge Taylor S, Tchernev VT, Mill Alsobrook JP, Gerlach V, Ec Macdougall J, Malyankar U, Gunther E, Stone DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to coding sequences for NOV proteins (ABN85378-ABN85403 and ABB98401-ABB98424). The NOV proteins and coding sequences are useful for treating or preventing NOV-associated disorders or in the manufacture of a medicament for treating the disorders, such as cancer, heart disease, inflammation, autoimmune disorders, allergies, blood disorders, AIDS, diabetes, obesity, asthma, IgA nephropathy, cirrhosis, arthritis, Alzheimer's disease, infections (e.g. bacterial, viral, parasitic), stroke, muscular dystrophy, epilepsy, and other wasting disorders associated with chronic diseases. The present sequence is a PCR primer for a NOV coding sequence, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JAN-2001;
28-FEB-2001;
09-MAR-2001;
                                                                                                 Human; allele-specific base detection; base-specific detection primer; allele-AS; high throughput; single nucleotide mutation detection; genetic variation;
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NOVX polypeptides and enc
treating NOVX-associated
Alzheimer's disease, and
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                                                                                                                                                                                            14-MAY-2003
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                                                                                                                                                                                                                                                                                                                                  3683
                                                             sapiens.
                                                                                                                                                                  genomic
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                                                                                                 llele-specific base detection; primer extension reaction; clific detection primer; allele-specific primer extension assay; throughput; single nucleotide polymorphism; SNP analysis; detection; genetic variation; allele-specific extension; prime
                                                                                                                                                                                                                                            standard;
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                                                                                                                                                                                                                                                                                                            GCCAGAAAGGCAACTATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page
                                                                                                                                                                                                                                                                                                                                                                                                              B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; 2001US-0260417P.
; 2001US-0260831P.
; 2001US-0272338P.
; 2001US-0274876P.
; 2001US-0284704P.
                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                  DNA mthfr SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            316;
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EA, Burgess CE, Gu
v VT, Miller CE, Gu
lach V, Edinger S, F
Yankar U, Millet I,
                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                        0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                encoding polynucleotides, useful for preventing ted disorders e.g. cancer, inflammation, or and in chromosome mapping, tissue typing or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                                                                  primer
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Pred. No. 2.1e
0; Mismatches
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Guo X, Boldog FL, Grosse
3, Rothenberg ME, Ellerman
I, Peyman J, Smithson G;
                                                                                                                                                                                                                                                                                                                                                                                                              ₽;
                                                                                                                                                                                                                                                                                                                                                                                                              0 Other
                                                                                                                                                                                                                                                                                                                                                                         1e+03;
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CC determined position in a nucleic acid molecule. The method comprises
CC performing primer extension reactions using base-specific detection
CC primers, each being specific for a particular base at the predetermined
CC position. The allele-specific (AS) primer extension assay method of the
CC invention is useful for detecting an allele-specific base at a pre-
CC determined position in a nucleic acid molecule, for high throughput
CC single nucleotide polymorphism (SNP) analysis, and for detecting
CC mutations and genetic variations. The new method solves the deficiencies
CC of previous methods by providing a method of allele-specific extension
CC that allows accurate discrimination between matched and mismatched
CC configurations, as well as reducing or eliminating false positive results
CC observed in prior art. The use of two allele-specific primers increases
CC the sensitivity by a factor of two because signals of two extensions are
CC obtained. The present sequence represents a primer used in the examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Sim
Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting a base at a pre-determined position in a nucleic acid molecule, comprises performing primer extension reactions using base-specific detection primers in the presence of a nucleotide-degrading enzyme.
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23-FEB-2001; 2001US-00791190.
07-FEB-2002; 2002US-00071926.
                                                                                                                                                                          Human; allele-specific base detection; primer extension reaction; base-specific detection primer; allele-specific primer extension a AS; high throughput; single nucleotide polymorphism; SNP analysis; mutation detection; genetic variation; allele-specific extension;
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23-FEB-2001; 2001GB-00004560 23-FEB-2001; 2001US-00791190
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                                          22-FEB-2002; 2002WO-GB000794
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                                                                                                                                 sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                        standard;
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                                                                                                                                                                                                                                                  primer
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RESULT 3144
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                          Human; cystic fibrosis; Tay-sachs; familial hypercholesterolaemia; FH; fragile X syndrome; haemophilia A; diabetes; cystinuria; tyrosinaemia; urea cycle disorder; hereditary fructose intolerance; Baker's disease; Wilson's disease; alcaptonuria; adult polycystic kidney disease; mCAD; Huntington's disease; myotonic dystrophy; retinitis pigmentosa; cancer; Gauchers disease; Canavan's disease; galactoseamia; thrombocytopaenia; thalassaemia; sickle cell disease; phenylketonuria; Marfan's syndrome; haemoglobinopathy; Bloom syndrome; Neimann Pick's disease; PCR; primer; galactose-1-phosphate uridyl transferase; GALT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          determined position in a nucleic acid molecule, for high throughput single nucleotide polymorphism (SNP) analysis, and for detecting mutations and genetic variations. The new method solves the deficiencies of previous methods by providing a method of allele-specific extension that allows accurate discrimination between matched and mismatched configurations, as well as reducing or eliminating false positive results observed in prior art. The use of two allele-specific primers increases the sensitivity by a factor of two because signals of two extensions are obtained. The present sequence represents a primer used in the examples
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08-MAY-2001; 2001US-00851501
                                                           06-MAY-2002; 2002WO-US014562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human GALT 8 specific PCR primer #2.
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Pred. No. 2.
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CC for identifying the presence or absence of a mutation or polymorphism in CC a subject, or the presence or absence of several genetic markers in a CC subject for diagnosing genetic diseases, e.g. cystic fibrosis, Tay-sachs, CC familial hypercholesterolaemia (FH), thalassaemia, sickle cell disease, CC phenylketonuria, galactosaemia, fragile x syndrome, haemophilia A, CC myotonic dystrophy, medium-chain acyl CoA dehydrogenase, maturity onset CC diabetes, cystinuria, methylmolonic acidaemia, urea cycle disorders, CC hereditary fructose intolerance, hereditary haemachromatosis, neonatal CC thrombocytopaenia, Gauchers disease, tyrosinaemia, Wilson's disease, CC alcaptonuria, hypolactasia, Baker's disease, argininaemia adenomatous CC polyposis coli (APC), adult polycystic kidney disease, Duchenne muscular CC dystrophy, alpha-1-antitrypsin deficiency, hereditary non-polyposis CC ecceal cancer, Huntington's disease, neurofibromatosis, Marian's CC syndrome, osteogenesis imperfecta, retinoblastoma, Freidrich's ataxia, CC haemoglobinopathies, MCAD, Canavan's disease, heber's hereditary oppic C neuropathy, retinitis pigmentosa, Bloom syndrome, Fanconi's anaemia, or Neimann Pick's disease. The present sequence is human galactose-1-CC phosphate uridyl transferase (GALT) specific PCR primer used to
Matches
                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AMBR-) AMBRY GENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method for identifying the presence or of a mutation or polymorphism in a plurality of genes. The method for identifying the presence or absence of a mutation or polymorph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying the presence or absence of a mutation or polymorphism in a subject, useful for diagnosing genetic diseases, comprises generating extension products and analyzing the melting behavior of the mixed DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunlop CLM,
                                                                                           Sequence 21
                                                                                                                                      illustrate the method
                       Local
l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 45; 49pp; English
                                                                                      BP; 5 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
Conservative
                     88.9%;
                                           0.2%;
                                                                                                                                      of the invention
0
                                             Score 14.8;
                          Pred.
  Mismatches
                          No.
                     2.1e+03;
                                             DB 1; Length
Indels
0,
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         absence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA stability; RNA expression; RNA synthesis; antisense; conzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme, G-Cleaver ribozyme; decoy molecule; aptamer; amberzyme, G-Cleaver ribozyme; tregion; viral replication; HBV reverse transcriptase; Enhancer I region; viral replication;
                                                                                                                                                                  17-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACD50312 standard; RNA;
                                                                                                                                                                                                                                                                                WO200281494-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               degenerative; disease state; HBV infection; HCV infection; cirrhosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HBV ribozyme substrate sequence #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        597 CTCCATCAAGTGGCTAGC 614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           failure; hepatocellular carcinoma; l
ide; antiinflammatory; substrate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hepatotropic;
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26-MAR-2002; 2002WO-US009187

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SXXXXXXXXXXXXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 3146
ACA60925
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to nucleic acid molecules which modulate CC the synthesis, expression and/or stability of Hepatitis C virus (HCV) or C the control of the con
                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Blatt L, .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-2001;
08-JUN-2001;
08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
05-DEC-2001;
                                                                                                                                                                11-AUG-2003
                                  Caenorhabditis elegans
                                                                                                                            Caenorhabditis elegans SUR-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel compound useful for hepatocellular carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (LEEP/)
(DRAP/)
                                                                                                                                                                                                                                     ACA60925 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MORR/)
                                                                                                                                                                                                                                                                                                                                                                   2904 TECCTTETTTCCTTETAT 2921
                                                                    8; Ras; PCR; cancer;
proliferation; cell
                                                                                                                                                                                                                                                                                                                                   2 ucacuucuuuccuucuau 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIBOZYME PHARM INC.
BLATT L.
MACEJAK D.
                                                                                                                                                                                                                                                                                                                                                                                                        6;
                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LEE P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001US-00877478.
2001US-0296876P.
2001US-0335059P.
2001US-0337055P.
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001US-00817879
                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 387pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 C; 1 G; 0 T; 11 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J,
                                                                                                                                                                                                                                         21
                                                                    Ras signalling; cell differentiation; growth; primer; Ras suppressor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating cirrhosis, liver failure, or condition associated with hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                      10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8;
Pred. No. 2.
                                                                                                                          exon 1 amplification primer
                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                         2.1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                        <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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ACC85521/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated nucleotide sequence encoding a suppressor of ras-8 (SUR-8) protein. Ras proteins are membrane bour GTpases. The nucleotide sequences are useful as targets for screeni drugs e.g. anticancer drugs, that can alter Ras signalling and the physiological effects of Ras on cell growth, differentiation and proliferation. The present sequence represents the Caenorhabditis e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleotide sequences encoding a suppressor of ras-8 protein, useful as targets for screening anticancer drugs that can alter Ras signaling, and the physiological effects of Ras on cell growth, differentiation an
                    New G-protein coupled receptor (GPCR) polypeptides and nucleic acids, useful for treating cancer, e.g. breast or prostate cancer.
                                                                                                                                                                                                                                        primer;
                                                                                                                                                                                                                                                   Human; G-protein coupled receptor; BCA-GPCR-1; BCA-GPCR-2; BCA-GPCR-3;
BCA-GPCR-4; breast cancer; cytostatic; cancer; prostate cancer; PCR;
                                                                                                                                                                                                                                                                                                            29-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proliferation.
                                                                                                                       20-DEC-2001; 2001US-00028521.
                                                                                                                                               18-DEC-2002;
                                                                                                                                                                       03-JUL-2003
                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                     Human G-protein coupled receptor PCR primer
                                                                                                                                                                                                                                                                                                                                  ACC85521;
                                                                                                                                                                                                                                                                                                                                                         ACC85521 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Han M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYTE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-JAN-2003.
                                                                             Powers
                                                                                                    (TULA-)
                                                                                                                                                                                            WO2003054542-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                              6389 AAAGCTCCTAATGCCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2003-415431/39
                                                      2003-569288/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exon 1 amplification primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7; Col 41; 67pp; English
                                                                            Ś
                                                                                                                                                                                                                                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                          88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UNIV TECHNOLOGY CORP
                                                                                                   TULARIK INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                  AAATCTCATAATGCCCAC
                                                                            Yang
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                 2002WO-US041112
                                                                                                                                                                                                                                                                                                            (first
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                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 6 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                             Cutler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                 21
                                                                                                                                                                                                                                                                                                                                                                                                                                       6406
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8; D:
Pred. No. 2.1e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding a
mbrane bound
                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and
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Example 1; Page 59; 77pp; English

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XSSSSSXX
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ADA09668
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Best Local S
Matches 16
The invention relates to an isolated nucleic acid molecule encoding a protein-coupled receptor HGPRBNY26. The nucleotide sequence comprises sequential nucleotide deletions from either the C-terminus or the N-terminus. Also included are an isolated polypeptide encoded by the nucleic acid, a recombinant vector comprising the nucleic acid, making recombinant host cell comprising the nucleic acid, diagnosing a
                                                                                                                                                                                                                                                                                                                                             07-MAR-2001;
27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention provides the protein and coding sequences of four novel human G-protein coupled receptors (GPCR) which are amplified in breast cancers. The sequences are useful in the treatment of cancers, including breast and prostate cancers. The present sequence is a PCR primer used to isolate a coding sequence for a GPCR of the invention
                                                                                                                                                                                                                                                                                       (FEDE/)
(RAMA/)
(MINT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cushing's syndrome; hyperprolactinaemia; haemochromatosis; congenital adrenal hyperplasia; follicle stimulating hormone deficiency;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            epididymitis; genital wart; germinal cell aplasia of the testis; cryptorchidism; varicocele; immotile cilia syndrome; viral orchitis; premature puberty; incomplete puberty; Kallman syndrome;
                                                                                                                                                                                                                                                                                                                                                                                      07-MAR-2002; 2002US-00092771.
                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003064381-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     testicular cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA09668
                                                                                                    Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        testicular cancer; choriocarcinoma; nonseminoma; seminoma; seminoma; spermatogenesis; infertility; Klinefelter's syndrome; XX m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-NOV-2003
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                                                                                                                                                                                                     2003-555589/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reproductive condition; amine disorder; testicular disorder;
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                                                                                                                                                                                                                                                        MINTIER G A.
CACACE A.
BARBER L E.
                                                                                                                                                                                                                                                                                                    FEDER J N.
RAMANATHAN C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein-coupled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTGGAGTTACTTTCATAG 3911
                                                                                                                                         G-protein coupled receptor HGPRBMY26 polypeptides and nucleic eful for preventing, treating or ameliorating e.g. testicular choriocarcinoma, infertility, viral orchitis, or Cushing's
                                                                                                  Page
                                                                                                                                                                                                                              Ramanathan CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HGPRBMY26; G protein-coupled receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 9 A; 5 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                             2001US-0273963P.
2001US-0278927P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                  78; 149pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor HGPRBMY26 Taqman
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                                                                                                                                                                                                                              Mintier GA,
                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.1e+03
                                                                                                                                                                                                                             Cacace A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR
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                                                                                                                                                                                                                              Barber LE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        XX male;
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                making
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC a member consisting of NFAT/CRE or NFAT G alpha 15, all undefined), and CC screening for candidate compounds capable of modulating activity of the CC HGPRBMY26 polypeptide. The HGPRBMY26 polypeptides, polynucleotides, compounds or pharmaceutical preparations comprising HGPRBMY26 are useful for preventing, treating or ameliorating a male reproductive condition; can amine disorder or a condition where 6-protein coupled receptors are (in)directly involved in disease progression, a testicular disorder, testicular cancer, choricocarcinoma, nonseminoma, seminoma, serinoma, sprintal warts, germinal cell aplasia of the testis, cryptorchidism, varicoccele, immotile cilia syndrome, viral orchitis, premature puberty; incomplete puberty, Kallman syndrome, Cushing's syndrome, hyperprolactinaemia, haemochromatosis, congenital adrenal C hyperplasia, follicle stimulating hormone (FSH) deficiency and C cranulmatons disease The present accounter in the content of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC16552 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression interference; expression inhibition; target gene; short interfering double stranded RNA; cytostatic; gene ther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          testicular tissue of a subject, identifying a compound that modulates the biological activity of a human G-protein coupled receptor HGPRBMY26 (and a member consisting of NFAT/CRE or NFAT G alpha 15, all undefined), and
                                                                                                                                                                                                                    Caplen NJ,
Kallioniemi
                                                                                                                                                                                                                                                                                                        (USSH )
(CARN-)
                                                                                                                                                                                                                                                                                                                                                                      30-JUL-2001;
08-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JUL-2002; 2002WO-US024226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003012052-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Short interfering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC16552;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           granulomatous disease. The present sequence is a reverse transcriptase (RT)-PCR probe used to determine the tissue expression profile of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pathological condition or a susceptibility to a pathological condition testicular tissue of a subject, identifying a compound that modulates t
                                                                                                                                                                           WPI; 2003-248169/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
                                                                                                                                                                                                                                                                               US DEPT HEALTH & HUMAN SERVICES. CARNEGIE INST WASHINGTON. ITI IMPERIAL COLLEGE INNOVATIONS LTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCCCACGGACTCCCAGCG
                                                                                                                                                                                                               Morgan RA, Fi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                      2001US-0308640P.
2002US-0370970P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double-stranded RNA oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 C; 4
                                                                                                                                                                                                               Fire A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                    INNOVATIONS LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8;
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                                                                                                                                                                                                                 Parrish S
Alton EW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.1e+03;
                                                                                                                                                                                                                 S, Mousses S;
, Griesenbach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NO:277.
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length of 0-nucleotide to 5-nucle genetic and/or therapeutic tools of a target gene.

for interfering

a 3' or 5' overhang each strand, useful fering or inhibiting

having expression

New RNA comprising double stranded RNA and length of 0-nucleotide to 5-nucleotides on

Claim 71; SEQ ID

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277; 176pp; English

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RESULT 3150
ADD14587
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cc inhibition of expression of a target gene, where (I) comprises double contained RNA of 15-40 nucleotides in length and a 3' or 5' overhang cc having a length of 0-nucleotide to 5-nucleotides on each strand, where cc having a length of 0-nucleotide to 5-nucleotides on each strand, where cc having a length of 0-nucleotide to 5-nucleotides on each strand, where cc portion of a mRNA or transcript of the target gene. Also described: (1) cc interfering with or inhibiting the expression of a target gene in a cell cc by exposing the cell to an amount of (I); (2) a gene silencing array comprising a substantially flat substrate, and addressably arrayed cc different double-stranded RNAs; (3) an array-based method of assessing a cc phenotypic effect of a double-stranded RNA on a target gene; (4) cc (5) selecting an optimised sequence of a double-stranded RNA for condition; (5) selecting an optimised sequence of a double-stranded RNA for condition; (6) a short double-stranded RNA effective for interfering with or condition expression of a target gene in a cell; cc inhibiting expression of a target gene comprising any of 311 20-78 cc activity, and can be used in gene therapy. The RNAs are useful as reverse genetic and/or therapeutic tools for interfering or inhibiting expression of a target gene in a cell; co fa target gene (see and constant con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  predictor set; protein tyrosine kinase activity modulator; protein tyrosine kinase; cytos gene therapy; drug sensitivity; genetic profile; cancer; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human src biomarker reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD14587 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes an RNA (I) used for the interference or inhibition of expression of a target gene, where (I) comprises double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                              New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein
                                                                                                                                                                                                                                                    WPI; 2003-636735/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-JAN-2003; 2003WO-US001981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003062395-A2
                                                                                                                                                                                                                                                                                                                                                                                                                  18-JAN-2002; 2002US-0350061P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                            (BRIM ) BRISTOL-MYERS SQUIBB CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4926 GACTGTTGAGTAACTCCT 4943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    target gene. They are useful for treating proliferative diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GACTGCTGAGGAACTCCT 3
                                                                                                                                 kinase pathways.
                                                                                                                                                                                                                                                                                                     Fairchild CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 4 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6 C; 5
                                                                                                                                                                                                                                                                                                        Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; D
Pred. No. 2.1e
0; Mismatches
                                                                                                                                                                                                                                                                                                     FΥ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer SEQ ID NO:776
                                                                                                                                                                                                                                                                                                        Shaw
                                                                                                                                                                                                                                                                                                        סי
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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The present invention describes a predictor set comprising a plurality polynucleotides or polypeptides whose expression pattern is predictive

요 요

SEQ

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776;

139pp; English

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the response of cells to treatment with a compound that modulates protein cyrosine kinase activity or members of the protein tyrosine kinase compound is capable of compathway. Also described: (1) predicting whether a compound is capable of compathway. Also described: (1) predicting whether a compound is capable of compathway. Also described: (2) predicting whether a compound is capable of compathway. Also described: (2) predicting obtaining a sample of cells, (2) determining a sample of cells, and (2) dentifying polynucleotides and polypeptides whose expression levels (2) correlate with compound sensitivity or resistance of cells associated with a disease state, comprising subjecting the plurality of cell lines to one or more compounds, analysing the expression pattern of a microarray of polynucleotides or polypeptides, and selecting (2) polynucleotides or polypeptides, and selecting (2) polynucleotides or polypeptides that predict the sensitivity or cells associated with a disease state by using the compounds that interact with protein tyrosine therapy. The polynucleotides and polypeptides are useful in predicting the cativity of compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways. These may be used in determining drug sensitivity in patients to allow the development of individualized cancer) based on patient response at a molecular level. The present of cancer) based on patient response at a molecular level. The present invention.
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RESULT 3151
ADD14405/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy; drug sensitivity; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                             predictor set; protein tyrosine kinase activity modulator; protein tyrosine kinase pathway; protein tyrosine kinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD14405 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 2 A; 7 C; 4 G; 8 T; 0 U; 0 Other;
                                New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein
                                                                                                                                                                                                                                                                             31-JUL-2003.
                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human src biomarker reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD14405;
                                                                                                                                                               (BRIM ) BRISTOL-MYERS SQUIBB CO
                                                                                                                                                                                                    18-JAN-2002; 2002US-0350061P
                                                                                                                                                                                                                                         17-JAN-2003; 2003WO-US001981.
                                                                                                                                                                                                                                                                                                                  WO2003062395-A2
                                                                                                                                                                                                                                                                                                                                                                                                           primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4219 TCCTTCCTCTGTGCAGAT 4236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 TCCGTCTTCTGTGCAGAT 19
                                                                                                                           Fairchild
                                                                                                                                                                                                                                                                                                                                                                                                            88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                             8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21
                                                                                                                               Lee FY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer SEQ ID NO:594.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                   tyrosine kinase; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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The present invention describes a predictor set comprising a plurality of coplynucleotides or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein cyrosine kinase activity or members of the protein tyrosine kinase compound that modulates protein tyrosine kinase expression of the markers of the protein tyrosine kinase creativity of cells, comprising obtaining a sample of cells, compound is capable of modulating the activity of cells, comprising obtaining a sample of cells, compound that mind a sample of cells, compound that the compound is capable of certaining whether the cells, comprising obtaining a sample of cells, compound that in the compound that the compound that the compound sensitivity of cell lines for correlate with compound sensitivity or resistance of cells associated with a disease state, comprising subjecting the plurality of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines to one or more compounds, analysing the expression pattern of cell lines or polypeptides, and selecting the compound that predict the sensitivity or resistance of cells associated with a disease state by using the capacity of compounds that interact with protein tyrosine kinases and/or genetic profiles which aid in treating diseases and disorders (e.g. cancer) based on patient response at a molecular level. The present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 3152
ADE27642/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 16
 11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
20-SEP-2002;
15-JAN-2003;
                                                                                                                                                                                                                                                                                                                            stearcyl-CoA desaturase; RNA interference; ancrectic; antidiab antiarteriosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening;
                                                                                                                                                                                                                                                                                                                 genetic
                                                                                                                                                                                                                                                                                                                                                                                     short interfering nucleic acid; siNA; downregulation; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                         Stearoyl-CoA desaturase siNA oligonucleotide SEQ ID NO:586.
                                                                                                                               20-FEB-2002;
                                                                                                                                                                   13-FEB-2003;
                                                                                                                                                                                                                                          WO2003070885-A2
                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE27642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                 engineering;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCTGGTGGGACAGTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCAGGTGGGAAAGTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP;
2002US-0358580P.
2002US-0386782P.
2002US-0406784P.
2002US-0406784P.
2002US-0409293P.
2002US-04029304P.
2003US-044012304P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                   2003WO-US004317.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 C; 1 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   594; 139pp; English
                                                                                                                                                                                                                                                                                                                 pharmacogenomic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1;
Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>, .</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                      antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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밁 5

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Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                that downregulates expression of the SCD (stearcy)-CoA desaturase) gene by RNA interference. Also described: (1) modulating expression of SCD genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA, and (4) vectors that express siNA. SCD inhibiting siNAs have anorectic, antidiabetic, antiarteriosclerotic, cytostatic and virucide activities. The siNAs can be used to modulate expression of SCD genes, in cells, tissue explants or organisms, e.g. for treating obsestly; diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; atherosca function and validation; genetic engineering; pharmacogenomics;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM
                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                    studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid, useful e.g. diagnosis of obesity or diabetes, downregulates stearoyl-CoA desaturase gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention describes a short interfering nucleic acid (downregulates expression of the SCD (stearcyl-CoA desaturase)
7232 TCCCTCTCAAGTCCAGCA 7249
                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; SEQ ID
                                                                                                                                                                     BP;
                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beigelman L,
                                                                                                                                                                     S
                                                                                                                                                                  A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO 586; 139pp;
                                                                                                                                                                  2 C; 8 G;
                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thompson
                                                        0
                                                                               Score 14.8; DB 1
Pred. No. 2.1e+03
                                                                                                                                                                  2 T; 4 U;
                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                     0 Other;
                                                                                                             DB 1;
                                                        2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for treatment and expression of the
                                                                                                          Length
                                                        Indels
                                                        0
                                                        Gaps
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                                                        0
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RESULT 3153
ADE47931/c
                                   Homo
                                                                     primer;
                                                                                      nootropic; antiparkinsonian;
                                                                                                     human; cardiant; antiarteriosclerotic; hypotensive; immunosuppressive; dermatological; anorectic; cytostatic; antidiabetic; haemostatic; anti-HIV; antiastimatic; antibacterial; virucide; neuroprotective;
                                                                                                                                                                               Human NOVX reverse PCR
                                                                                                                                                                                                                  29-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                    931/c
ADE47931 standard; DNA;
WO2003076642-A2
                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                   18
                                                                                                                                                                               primer
                                                                                                                                                                                                                                                                                            21 BP
                                                                                                                                                                                                                                                                                                                                                                                   Ļ
                                                                                      antilipaemic; gene therapy;
                                                                                                                                                                               SEQ
                                                                                                                                                                               ij
                                                                                                                                                                                 NO:293
                                                                                        PCR;
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02-AUG-2001; 03-AUG-2001; 08-AUG-2001; 09-AUG-2001; 13-AUG-2001; 14-AUG-2001; 17-AUG-2001; 17-AUG-2001;

2001US-0309501P.
2001US-0310291P.
2001US-0310951P.
2001US-0311292P.
2001US-0311292P.
2001US-0312156P.
2001US-0313150P.

02-AUG-2002; 18-SEP-2003.

2002WO-US024459

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                                   ક
                                                                                                                                                                                                          neuroprotective, nootropic, antiparkinsonian, and antilipaemic activity.

Characteristic protective, nootropic, antiparkinsonian, and antilipaemic activity.

Characteristic protective, nootropic, antiparkinsonian, and antilipaemic activity.

Characteristic protection as a vaccine. A polypeptide of the invention is considered with the manufacture of a medicament for treating a syndrome associated with the polypeptide. These manufacts are apathology consisted with the polypeptide. These manufacts such as cardiomyopathy, creating or preventing NOVY-associated disorders such as cardiomyopathy, catheroscierosis, hypertension, scleroderma, obesity, cancer, diabetes, canternoscial, graft-versus-host disease, AIDS, asthma, Crohn's disease, comultiple sclerosis, infections, anorexia, cancer-associated cachexia, neurodegenerative disorders (e.g. Alzheimer's disease or Parkinson's concerdes associated with chronic diseases. The nucleic acids are also cused as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine, and pharmacogenomics. The polypeptides are also concerdes associated with chromic diseases.
                                                                      Matches
                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-AUG-2001;
21-SEP-2001;
03-DEC-2001;
05-PEB-2002;
05-MAR-2002;
19-APR-2002;
15-MAY-2002;
15-MAY-2002;
28-MAY-2002;
28-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-AUG-2001;
23-AUG-2001;
28-AUG-2001;
29-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pena CEA, Sh
Boldog FL, P.
Vernet CAM,
Burgess CE,
Chaudhuri A,
Glot L, Ool
                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunosuppressive, dermatological, anorectic, cytostatic, antihaemostatic, anti-HIV, antiasthmatic, antibacterial, virucide, neuroprotective, nootropic, antiparkinsonian, and antilipaemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, diabetes, atherosclerosis, asthma or AIDS, and in chromosome mapping, tissue typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel (NOVX) human polypeptide. A polypeptide of the invention has cardiant, antiarteriosclerotic, hypotensive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-779062/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Giot L, Oc
Taupier RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-AUG-2002;
                                                                                                                                                                                               preventive medicine, and useful as vaccines. The
                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP
                                   7384
 19
                                                                      1 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BD, Patturajan M, Kekuda R, Miller CE, Rieger DK;
Shimkets RA, Li L, Berghs C, Zhong M, Casman SJ, Voss EZ;
L, Padigaru M, Smithson G, Shenoy SG, Ji W, Gorman L;
AM, Leite MW, Guo X, Anderson DW, Spytek KA, Gerlach VL;
CE, Khramtsov NV, Ort T, Ellerman K, Rastelli L, Agee ML;
LA, Chant JS, Dipippo VA, Eddinger SR, Elsen A, Gangolli EA;
Ooi CE, Rothenberg ME, Spaderna SK, Hjalt T, Liu X;
                                                                                                                                             21 BP; 7
                                 TGTACAGTTCCTTCTGAA 7401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 293; 562pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001US-0314031P
2001US-0315405P
2001US-0315803P
2001US-0315803P
2001US-0323936P
2001US-0338078P
2002US-0354655P
2002US-0351764P
2002US-0373825P
2002US-036071P
2002US-0380971P
2002US-0380980P
2002US-0380980P
2002US-0380980P
2002US-0383761P
2002US-0383887P
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Catterton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002US-00210130
                                                                                                                                             ?
                                                                                   0.2%;
                                                                                                                                             3 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E
                                                                                                                                                                                                 present sequence represents
                                                                      <u>.</u>
                                                                                                                                             G; 3 T; 0 U; 0 Other;
                                                                                       Score 14.8;
Pred. No. 2.
                                                                      Mismatches
                                                                                       2.1e+03
                                                                                                         DB 1;
                                                                                                         Length 21;
                                                                        Indels
                                                                                                                                                                                                 a PCR primer used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antidiabetic,
                                                                      0;
                                                                      Gaps
                                                                      0,
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RESULT 3154
AAT78996
ID AAT78996 standard; DNA; 22

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RESULT 3155
AAT94992/c
ID AAT94992/c
XX AAT9499
XC AAT9499
XY O2-APR-
XX Primer
XX Human |
KW locus |
XX Synthet
OS Synthet
OS Homo si
XX W09731:
XX PD 28-AUG-
XX PR 20-FEB-
XX PR 20-FEB-
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This oligonucleotide comprises the 5' acceptor site of intron 1 of the human Huntington's disease (HD) gene. The splice site sequences for the first 5 exons of the mouse HD gene (see AAT78974) and the human HD gene were compared (see AAT7895-T79002). Targeted disruption of the murine HD gene, e.g. at exon 5, can be used to examine the function of the HD gene and its role in development. Transgenic mice can be used as models of HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 BP; 2 A; 3 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse Huntington's Disease gene - useful for generating transgenic mice as a model of Huntington's Disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Huntington's disease; animal model; transgenic animal; human; therapy; drug screening; Hdh gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 60; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-298677/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hayden M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT78996;
                     20-FEB-1996;
                                                        20-FEB-1996;
                                                                                                                                    WO9731126-A1
                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                 Primer 6 for sequencing of human leukocyte antigen class I genes.
                                                                                                                                                                                                                                                                                                                      02-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                AAT94992 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYBR-) UNIV BRITISH COLUMBIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4463 CITITITITITITITI
                                                                                                                                                                       sapiens
                                                                                                                                                                                                                             leukocyte antigen-C class I gene; HLA-C; specific nucleic acid amplification; HLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Huntington's disease gene intron 1 3' acceptor site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lin B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95US-00457273.
                       96WO-US002408
                                                          96WO-US002408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96CA-02178022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nasir J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                               exon 1;
typing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ,
,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 3156
AAT95005/c
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primers were used for locus specific nucleic acid amplification for HLA typing. Typing HLA-A, -B or -C class 1 genes comprises providing a sample DNA containing a HLA-A, -B or -C class 1 gene having a 1st and 2nd exon and a target sequence, contacting the sample DNA with an amplification primer including sequence complementary to sequence located in exon 1 of the HLA-A, -B or -C gene, and a second amplification primer sequence complementary to sequence located in exon 5 of the HLA-A, -B or -C gene. The PCR product is sequenced using the above primers and the determined DNA sequence compared with the DNA sequences of known HLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human
locus
The present sequencing primer was used leukocyte antigen (HLA) class I genes.
                                                                                                                                                                                                                                                                                                                                                                                                                         WO9731126-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing primers AAT94987-92 were used to sequence PCR amplified hum leukocyte antigen (HLA) class I genes. The primers are designed to hybridise to exon-intron borders of exons 2, 3 and 4 of the HLA genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amplification and sequencing primers useful for locus specific nucleic aci
                                                                                                    Amplification and sequencing useful for locus specific nuc
                                                                                                                                                                    WPI; 1997-435175/40.
                                                                                                                                                                                                                                                                                            20-FEB-1996;
                                                                                                                                                                                                                                                                                                                                      20-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT95005 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Johnston-Dow L,
                                                                                                                                                                                                                                                     (PEKE ) PERKIN-ELMER CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PEKE ) PERKIN-ELMER CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       leukocyte antigen-C class I gene; HLA-C; exon 1; exon 5; specific nucleic acid amplification; HLA typing; exon 4;
                                                             29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10; Page 57; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCTGCCGAAGCCCCTCAC 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCAGCCCAAGCCCCTCAC 959
                                                             Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequencing exon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                             62;
                                                                                                                                                                                                                                                                                            96WO-US002408
                                                                                                                                                                                                                                                                                                                                      96WO-US002408
                                                                                                                                                                                                           Chadwick RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chadwick RB,
                                                           105pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22
                                                             English
                                                                                                    ing primers
nucleic aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 sense strand of HLA class
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1;
Pred. No. 2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G; 3 T; 0
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                                                                                                    ers specific for HLA class 1 acid amplification for HLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ers specific for HLA class 1 genes acid amplification for HLA typing
The primer is designed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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                                                                                                                      1 genes
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the sense strand of exon 4, from the 5' exon-intron border. PCR primers were used for locus specific nucleic acid amplification for HLA typing. Typing HLA-A, -B or -C class 1 genes comprises providing a sample DNA containing a HLA-A, -B or -C class 1 gene having a lst and 2nd exon and target sequence, contacting the sample DNA with an amplification primer including sequence complementary to sequence located in exon 1 of the HLA-A, -B or -C gene, and a second amplification primer sequence complementary to sequence so of the HLA-A, -B or -C gene. The PCR product is sequenced using the above primers and the determined DNA sequence compared with the DNA sequences of known HLA types
                                        The specification describes termamyl-like alpha-amylase variants that have altered amino acid sequences to improve properties. The variants are produced by creating one or more of the following mutations in amino acid sequence of the parent termamyl-like alpha-amylase: T141, K142, F143, B144, F145, F146, G147, R148, G149, Q174, R181, G182, D183, G184, K185, A186, W189, S130, N195, H107, K108, G109, D166, W167, D168, Q169, S170, R171, Q172, F173, F267, W268, K269, W270, D271, L272, G273, A274, L275, K311, E346, K385, G456, N457, K458, P459, G460, T461, V462, T463. The variants can be used for washing and/or dishwashing. They can also be used in the production of sweeteners and ethanol from starch, and/or textile desizing, and in starch liquefaction and/or saccharification processes. The present PCR primer was used to construct the variants of
                                                                                                                                                                                                                                                                                                                             WPI; 1999-326987/27.
                                                                                                                                                                                                                                                                                                                                                            Borchert TV,
Kjaerulff S;
                                                                                                                                                                                                                                                                                                                                                                                                                                            30-OCT-1997;
14-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX59677 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                          Example 10; Page 58; 115pp; English
                                                                                                                                                                                                                                                                                           New Termamyl-like alpha-amylase variants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Termamyl-like; alpha-amylase; variant; washing; dishwashing;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           942 GCAGCCCAAGCCCCTCAC
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98DK-00000936.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 C; 11 G; 3
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                                                                                                                                                                                                                                                                                                                                                                           Andersen C, Nielsen BR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         textile desizing; starch liquefaction;
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Pred. No. 2.2e+03;
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RESULT 3158
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XX MAGE AAA6461
XX MAGE AAA6461
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YX WPI; 20
CC Complen
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PT Complen
PT Cancer
XX WPI; 20
CC Canciger
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CC Gescril
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RESULT 3159
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AC AAA9055
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Matches 16
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Serrano A, I
Boon-Falleur
                           11-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genes encode tumour rejection antigens which complex to human leukocyte antigens (HLAs), and provoke response by autologous, cytolytic T cells. The genes are located in region q28 of the X chromosome. The MAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primers AAA64618-19 were used in reverse transcription PCR reactions to assess the frequency of expression of MAGE-A10 gene. The specification describes MAGE-A5, MAGE-A8, MAGE-A9, MAGE-A10 and MAGE-A11. The MAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JAN-2001
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                                                                       AAA90556;
                                                                                                                   AAA90556
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 4; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Complementary polynucleotide of MAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAR-2000; 2000WO-US005346.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA64619;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polynucleotides are useful for diagnosis of cancer
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                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                     standard;
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                                                                                                                                                                                                                                         ACCTCCTCAGGGGTGCAG
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                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                assess frequency of expression of MAGE-Al0
                                                                                                                                                                                                                                                                                                                                                       0.2%;
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                                                                                                                                                                                                                                                                                                                                 Score 14.8; D
Pred. No. 2.2e
0; Mismatches
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Pred. No. 2.2e+03;
D; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 3160
AAA90562/c
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is a sequencing primer for Human Leukocyte Antigen (HLA) class I gene. The HLA class I genes are found on chromosome 6p21.3. HLA class I proteins are found on the surface of almost all nucleated cells and are involved in antigen presentation to immune system cells. This primer can be used to type HLA class I genes by carrying out PCR on a sample DNA, comprising HLA class I gene, and detecting the amplicon formed using a sequence-specific detection method e.g. DNA sequencing (using the present sequence). The present sequence is useful for discriminating among the HLA-A, HLA-B, and HLA-C genes and other related class I genes and pseudogenes. In addition, the present sequence is useful for organ transplantation studies, for the study of autoimmune disease and for the determination of susceptibility to infectious disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Typing HLA class I genes for organ transplantation, involves contacting the sample DNA containing HLA class I gene comprising two exons and a target sequence, with amplification primers and detecting the amplicon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Leukocyte Antigen class I; HLA-\lambda; antigen presentation; organ transplantation; autoimmune disease; sequencing primer; infectious disease susceptibility; chromosome 6p21.3; ss.
                                                                                                               Human Leukocyte Antigen class I; HLA-A; antigen presentation; HLA typing; organ transplantation; autoimmune disease; sequencing primer; infectious disease susceptibility; chromosome 6p21.3; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22
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                                                                                                                                                                              HLA class I
                                                                                                                                                                                                          11-JAN-2001
                                                                                                                                                                                                                                                                    AAA90562 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 40; Col 38; 60pp; English
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                                                      US6103465-A.
                                                                                                                                                                                                                                        AAA90562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PEKE ) PERKIN-ELMER CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-AUG-2000.
                                                                                    sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 3 A; 5 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95US-00390251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-00538666
                                                                                                                                                                            sequencing
                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                       BP
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                                                                                                                                                                            primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chadwick RB;
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1;
Pred. No. 2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                              #12.
                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                         2
                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
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03-OCT-1995;

95US-00538666.

15-AUG-2000.

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PROCESS OF A SPECIAL PROCESS O
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AAZ91959
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is a sequencing primer for Human Leukocyte Antigen (HHA) class I gene. The HLA class I genes are found on chromosome 6p21.3. HLA class I genes are found on chromosome 6p21.3. HLA class I proteins are found on the surface of almost all nucleated cells and are involved in antigen presentation to immune system cells. This primer can be used to type HLA class I genes by carrying out PCR on a sample DNA, comprising HLA class I gene, and detecting the amplicon formed using a sequence-specific detection method e.g. DNA sequencing (using the present sequence). The present sequence is useful for discriminating among the HLA-A, HLA-B, and HLA-C genes and other related class I genes and pseudogenes. In addition, the present sequence is useful for organ transplantation studies, for the study of autoimmune disease and for the determination of susceptibility to infectious disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mahogany gene; mouse; mg gene; regulatory defect; gene therapy; obesity; weight regulation; cell therapy; body weight disorder; cachexia; anorexia; hyperpigentation; increased metabolic rate disorder; hyperphagia; Antiobesity; antianorexic; anticachexic; PCR primer; ss.
New human and murine mahogany genes, useful, e.g. for diagnosis
                                                                                                                                                                                                                                                                   21-JUL-1998;
20-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ91959 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; Col 35; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Typing HLA class I genes for organ transplantation, involves contacting the sample DNA containing HLA class I gene comprising two exons and a target sequence, with amplification primers and detecting the amplicon.
                                                               WPI; 2000-195103/17.
                                                                                                                                                                                                                                           05-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                          21-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                   03-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200005373-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   942
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16; Conserv
                                                                                                                                                                                 MILLENIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GCAGCCCAAGCCCCTCAC 959
                                                                                                                  Nagle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Johnston-Dow L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 3 A; 5 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                       98US-0093630P.
98US-0104978P.
99US-00245041.
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                                                                                                                                                                                                                                                                                                                                                          99WO-US016484.
                                                                                                                        DĽ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    exon 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer CelegE117.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                          23-JUN-1999;
07-JUL-1999;
26-JUL-1999;
30-NOV-1999;
20-DEC-1999;
05-JAN-2000;
                                                                                                                                                                                                                                                                                                                                     proliferation; tumourigenesis; identification; cancer; PCK primer; hybridisation; probe; cytostatic; nootropic; neuroprotective; antiinflammatory; immunosuppressive; immunostimulant; antiangiogenic; leukaemia; lymphoid malignancy; neuronal disorder; glial disorder; estrocytal disorder; hypothalamic disorder; glandular disorder; macrophagal disorder; ppithelial disorder; stromal disorder; epithelial disorder; andiogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (cell-based) gene therapy. (II) are used to raise specific antibodies (Ab); to identify other (extra)cellular products involved in weight regulation, and to screen for agents that disrupt interaction between (II) and other macromolecules. The Ab are used to detect abnormal levels (or function) of (II) (for diagnosis, prognosis or monitoring of treatment); to evaluate (II)-expressing cells intended for cell therapy, and as therapeutic mg inhibitors. Cells that express the mg gene (or contain the mg polypeptide) are used to identify agents (A) that modulate mg activity. (A) are potentially useful for the treatment of body weight disorders, particularly obseity, cachexia or anorexia, or other conditions associated with the mg gene such as hyperpigmentation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention. The mahogany genes are used: (i) to produce recombinant mahogany (mg) proteins (II); (ii) as a source of antisense, ribozyme o triplex-forming therapeutics; (iii) as a source of diagnostic probes a primers for detecting expression of mg genes or mutations, regulatory defects, in this gene, or for isolation of related sequences; and (iv)
                                                                                                                                                                                                                                                       WO200053755-A2
                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO212 hybridisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence represents a PCR primer for a mahogany gene of the invention. The mahogany genes are used: (i) to produce recombina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Fig
                                                                                                                                            02-JUN-1999;
                                                                                                                                                           08-MAR-1999;
                                                                                                                                                                                          06-JAN-2000; 2000WO-US000376
                                                                                                                                                                                                                        14-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC58261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC58261 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hyperphagia and disorders that result in increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment of body weight disorders
                (GETH )
                                                                                                                                                                                                                                                                                                                  immunologic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7072 TGAATGCACTGAGTCCCT
                                                                                                                                                                                                                                                                                     sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                tumour;
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                GENENTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                               2000WO-US000219
                                                                                                                                                                                                                                                                                                                                   disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5; 188pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                         99US-0141037P.
99US-0143048P.
99US-0145698P.
99WO-US028313.
99WO-US030911.
                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis; neoplastic disease; neoplastic cell growth; tumourigenesis; identification; cancer; PCR primer;
                                                                                                                                                           99WO-US005028
                                                                                                                                            99WO-US012252
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                INC
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                                                                                                                                                                                                                                                                                                                    88.
                                                                                                                                                                                                                                                                                                                                 inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probe SEQ ID NO:80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7089
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Pred. No. 2.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.2e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
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RESULT 3163
ABA82014 standard; DNA; 22 BE
XX
AC ABA82014;
XX
DT 25-JAN-2002 (first entry)
XX
Mouse wound healing related E
XX

Muse wound healing related E
XX

Muse musculus.
XX
PN (CA2325226-A1.
XX
PN (CA2325226-A1.
XX
PD 17-MAY-2001.
XX
PD 17-NOV-1999; 99US-0172511P
PR 17-DEC-1999; 99US-0172511P
PR 20-JUN-2000; 2000DE-01030149
XX
PA (SWIT-) SWITCH BIOTECH AG.
XX
PA (SWIT-) SWITCH BIOTECH AG.
XX
Regenbogen J, Wolf E, Gopp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an isolated antibody that binds to one of the human PRO proteins designated PRO212, PRO290, PRO314, PRO535, PRO619, PRO1017, PRO809, PRO3107, PRO1005, PRO1005, PRO1005, PRO1025, PRO1030, PRO1097, PRO1107, PRO1111, PRO1153, PRO1105, PRO11087, PRO11097, PRO1107, PRO1111, PRO1151, PRO1153, PRO1184, PRO1187, PRO1181, PRO2139, PRO39, PRO834, PRO1117, PRO1151, PRO2094, PRO2145 OR PRO2198. PRO antagonists can be used to inhibit tumour cell growth. The CC PRO1281, PRO23, PRO39, PRO834, PRO1117, PRO1710, PRO2094, PRO2145 OR PRO2198. PRO antagonists can be used to inhibit tumour cell growth. The CC PRO polypeptides and nucleotides are useful in the treatment, diagnosis and prevention of cancer. The antibodies and other anti-tumour compounds cand prevention and/or activation of the amplified PRO genes. Exemplary CC conditions or disorders to be treated with such antibodies and other CC compounds include benign or malignant tumours (e.g., renal, liver, CC kidney, bladder, breast, gastric, ovarrian, colorectal, prostate, pancreatic, lung, vulva, thyroid, hepatic carcinomas, sarcomas, CC pancreatic, lung, vulva, thyroid, hepatic carcinomas, sarcomas, CC glioblastomas, and various head and neck tumours), leukaemias and CC lymphoid malignancies, other disorders such as neuronal, glial, CC stromal and blastocoelic disorders, and inflammatory, angiogenic and CC immunologic disorders. AAC58242 to AAC58366 represent PCR primers and CC myloridisation probes used in the isolation of the human PRO sequences.

CC polynucleotide and protein sequences given in the exemplification of the creation of the human PRO sequences.
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Best Local Similarity
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Watanabe CK,
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                                                                                                                                                                                                                                                                                                                                   healing related PCR primer SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                         99DE-01055349.
99US-0172511P.
2000DE-01030149.
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       Goppelt A,
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to the use of a polypeptide (ABB4454-ABB44601, ABB44606-ABB44623) or its variant or encoding nucleic acid (ABA81990-ABA81995-ABA82032) with vulnerary and/or dermatological activity for the diagnosis, prevention and treatment of skin disorders and treatment in wound healing or for the identification of pharmacologically active substances. The nucleic acids are useful in gene therapy. The present sequence is that of a PCR primer, useful to the invention. Note: The printed sequence listing for this specification was incomplete, terminating part way through SEQ ID NO 106. The remaining data was obtained from EPO data for an equivalent patent (EP1114862)
The present invention relates to a kit and method for the detection of gene polymorphisms of drug-metabolising enzyme genes. The kit contains a polymerase chain reaction (PCR) buffer solution containing DNA polymerase and NTP, a normal forward primer, a mutated forward primer, a reverse primer and a fluorescence-labelling probe. The method involves carrying out PCR on sample DNA, containing a drug-metabolising enzyme gene, together with PCR buffer, the normal forward primer, the reverse primer and the fluorescence-labelling probe (step A); and carrying out PCR on the sample DNA together with PCR buffer, the mutated forward primer, the
                                                                                                                                                                                    Detection
diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of novel polypeptide or its variant or nucleic acid encoding the polypeptide for diagnosing and/or preventing and/or treating skin disorders and/or treatment in wound healing or for identifying active
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                                                                                                                                                                                                                                  WPI; 2001-285409/30
                                                                                                                                                                                                                                                                                             19-MAR-1999;
06-MAY-1999;
                                                                                                                                                                                                                                                                                                                                         10-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism; drug-metabolising enzyme; PCR primer; CYP2C18i; ss.
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nilarity 88.9%;
Conservative (
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99JP-00125918
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present invention

reverse primer and the fluorescence-labelling probe (step B), and a step of comparing the result of step a with that of step b. The present sequence is a primer for human CYP2C181, which was used to illustrate the

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RESULT 3165
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                                                                                                                                                                                                                                 The present sequence represents a primer used to produce maize cDNA. specification describes plant androgenesis markers, which are derived from maize. Maize androgenesis markers are used to evaluate the androgenic potential of plants and derived cultures of microspores. I markers are also used to clone genes involved in androgenesis, and for genetic analysis to identify quantitative trait loci involved in androgenesis, particularly in cereals, specifically maize. Varieties capable of androgenesis are useful in plant breeding programs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New markers for plant androgenesis, useful for evaluating androgenic potential, cloning of androgenesis genes and genetic analysis.
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                                                                                                                                  Length 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                          0
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                                                                                                                                                                                                                                                                                                                                                                               derived
                                                                                                                                                                                                                                                                                                            and for
                                                                                       Gaps
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                                                                                          0
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RESULT 3166 AAH37409/c

밁 Ś

20

Matches

16;

Conservative

Mismatches

Indels

0

Gaps

0

2.2e+03

2411 CAGTGGACACCAACATCA

2428 0, Pred.

rocar

Similarity

88.9%;

RESULT 3167

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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP Ilanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid sample by CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC auged by one or more SNPs, Phenotypic traits include diseases e.g. CC osteogenesis imperfects and acute intermittent porphyria, Phenotypic CC traits also include symptoms of or susceptibility to multifactorial CC disease of which a component is or may be genetic such as autoimune CC disease of which a component is or may be genetic such as autoimune CC inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and content in the present semence represents a DCD primer specific.
Query Match
                                      Sequence
                                                                          paternity analysis. The prefor a human SNP containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 51; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNPE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH37409 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-290930/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNPE; genotyping;
Lesch-Nyhan syndro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH37409;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ORCH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORCHID BIOSCIENCES INC
                                        22 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0160096P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphism; SNP; single nucleotide primer extension;
agammaglobulinaemia; diabetes insipidus; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR
                                                                                              The present sequence
                                    4 C; 5 G; 10 T; 0 U; 0 Other;
  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Z
                                                                            DNA sequence
  Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID
                                                                                                represents a
  DB 1;
  Length
                                                                                                PCR primer specific
    22;
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AWH40497
III AWH4
XX
AC AWH4
AC AWH4
XX
AC AWH4
XX
AC AWH4
AC AWH4
XX
AC AWH4
AC AWH4
AC AWH4
XX
AC AWH4
AC A
                                                                                                                                                                                                                                                                                                                                Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide compilers of single nucleotide polymorphisms SNPs. The present invention concludes kits for determining the presence or absence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a component of the concludes of the invention. The PCR primers are used to amplify a component of the concludes are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The coligonucleotides are useful for determining the presence, absence or conditionable of a SNP and for genotyping nucleic acid sample by conditionable of a SNP and for genotyping nucleic acid sample by conditionable of a SNP and for genotyping nucleic acid samples, for e.g. to casees by association analysis the genotype of an individual or group of conditionable, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic trait suspected of being cagement by the presents include diseases e.g. consteogenesis imperfects and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial constants also include symptoms of or susceptibility to multifactorial confidences, including, rheumatoid arthritis, multiple sclerosis, concert, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and concert, nervous system diseases and infection by pathogenic concert, and susceptibility to multifactorial accordance of the present sequence represents a PCR primer specific concert, and susceptibility to constitutions and paternity analysis. The present sequence represents a PCR primer specific for a human SNP containing DNA sequence
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                                                          S
                                                                                                                                            Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 66; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP specific upper PCR primer SEQ ID 3293.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200129262-A2
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                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ORCH-) ORCHID BIOSCIENCES INC.
                                                                  4832 CAAAACAATCTATCCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sample.
   N
                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                        22
                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
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                                                                                                                                                                         0.2%;
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                                                                                                                                                                                                                                                                                    C; 1 G; 4 T; 0 U;
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                                                                  4849
   19
                                                                                                                                            <u>.</u>
                                                                                                                                        Score 14.8; D
Pred. No. 2.2e
0; Mismatches
                                                                                                                                                                             2.2e+03
                                                                                                                                                                                                              DB 1; Length 22;
                                                                                                                                                                                                                                                                                    0 Other;
                                                                                                                                        0
                                                                                                                                    Gaps
                                                                                                                                        0
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RESULT 3169 ABS59077/c ID ABS5907

ABS59077 standard; DNA; 22

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밁 ঠ

w

ATCAGCCAGAGATCTCTG

20 2982

2965 ACCAGCCAGAAATCTCTG

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CC check whether they are susceptible to toxins or resistant to certain CC check whether they are susceptible to toxins or resistant to certain CC damage, skin cancer and cancers associated with oxidative stress. The CC method allows the characterisation and mechanism of the GST1 deletion CC and identifies 18 kb homology regions flanking GSTT1 which are involved CC in the deletion event that produced the *0 allele. The method permits the CC unambiguous discrimination of all GST1 genotypes (*A), *0/0 (both CC homozygous), *A/0 (heterozygous)). The technique allows the reproducible CC simultaneous discrimination of all GST1 genotypes (*A), *0/0 (both CC enzyme activity in erythrocytes. The trimodular distribution of CC entrypes detected by these procedure correlated highly significant with CC phenotypes at high, intermediate and null- activity in homo- and CC indicate a gene dose effect. The present sequence is a PCR primer for CC amplifying a 3187bp fragment from the human GSTT1*0 allele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 3168
   Matches
                  Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                          the sequence upstream of the deletion area and the other is from the sequence downstream of the deletion area, and checking the corresponding DNA fragment produced in PCR. The method is useful for detecting presence of at least GSTT1*0 allele, for diagnostic testing of individuals to
                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a PCR assay for detecting presence of at least a single allele of deletion mutant GSTT1*0 (glutathione S-transferase theta allele) involves performing PCR with two primers, of which one is from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR assay to detect presence of single allele e.g., glutathione S-transferase thetaasteriskO allele, of deletion mutant involves performing PCR with primers derived from sequences upstream and downstream of deletion area.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP1130112-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; PCR primer; ss; GSTT1; Glutathione-S-transferase theta; skin cancer; GDTT1*0 allele; oxidative stress; genotyping; GST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human GSTT1*0 3187bp fragment PCR primer GST-TR9n
                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 17; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-591524/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sprenger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-FEB-2000; 2000EP-00103844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-FEB-2000; 2000EP-00103844.
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l Similarity
16; Conserv
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                                                                        22
                                                                        B₽;
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Schlagenhaufer R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                        σ
                                                                        A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                    0.2%;
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 0,
                    Score 14.
Pred. No.
   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ą
     2 14.8; L-
No. 2.2e+03;
                                     DB 1;
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                                   Length
   Indels
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   0
   Gaps
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The invention relates to novel isolated G-protein coupled receptor (GPCR) CC polypeptides and polynucleotides. The GPCR polypeptide, GPCR nucleic acid cand antibody are useful for treating, preventing or alleviating a GPCR-CC associated disorder or a pathological state in a subject, particularly a human. In particular, the disorder is cardiomyopathy, atherosclerosis, CC diabetes, or a disorder related to cell signal processing and metabolic control of the presence of or predisposition to a disease associated control of the presence of or predisposition to a disease associated control of the presence of or predisposition to a disease associated control of the presence of or predisposition to a disease associated control of the presence of or predisposition to a disease associated control of the presence of or predisposition to a disease associated control of the presence of or predisposition to a disease associated control of the presence of or predisposition or prophylactic acid and polypeptide are especially useful in therapeutic or prophylactic control of the protein associated with aberrant GPCR expression or control of the protein is useful in gene therapy for control of the protein is useful in gene therapy for control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the nucleic acids and control of the protein of the pro
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24-JAN-2001;
08-FEB-2001;
22-FEB-2001;
14-MAR-2001;
23-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Casman S
Gerlach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; G-protein coupled receptor; GPCR; cardiomyopathy; atherosclerosis diabetes; cell signal processing; metabolic pathway modulation; cancer; adenocarcinoma; prostate cancer; uterus cancer; asthma; immune response; neurodegenerative disorder; inflammatory disorder; Crohn's disease; multiple sclerosis; Albright hereditary osteodystrophy;
                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-JAN-2001;
10-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in gene therapy, atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New G
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23-APR-2001;
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21-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-599789/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein coupled receptor polypeptides and polynucleotides, useful ne therapy, particularly for treating or preventing cardiomyopathy, pacterosis, diabetes, multiple sclerosis, Crohn's disease or cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ballinger RA,
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PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURAGEN
                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vernet CA
Smithson
Ellerman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; 2001US-0259743P.
; 2001US-0261498P.
; 2001US-0261498P.
; 2001US-0263689P.
; 2001US-0271021P.
; 2001US-027946P.
; 2001US-027945P.
; 2001US-0278150P.
; 2001US-028571P.
; 2001US-028571P.
; 2001US-029327P.
; 2001US-0312902P.
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2000US-0257876P.
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                                                                                                                                                                                                                                                                                                                                                                                                                             685pp;
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M, Shenoy SG, Gusev V,
G, Stone DJ, Sciore P,
K, Gangolli EA, Millet
                                                                                                                                                                                                                                                                                                                                                                                                                             English
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Malyankar UM, Edi
Macdougall JR, Gu
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ABK33445/c
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Best Local S
Matches 16
                         Matches
                                    Query Match
Best Local
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                                                                                                                           The present invention relates to a method for detecting non-responders t anti-tumour necrosis factor (TNF) therapy. The method involves testing a individual for homozygosity for at least one single nucleotide polymorphism (SNP) in the gene coding for TNF receptor II, which is located on chromosome 1p36. Two novel SNPs, one in exon 2 (position 168 A/G) and one in exon 6 (position 587 T/G) which result in Lys56Lys and Met196Arg respectively, are also described. The method of the invention is useful for detecting non-responders to anti-TNF therapy such as inflixingly therapy, or therapy such as the interaction of the invention is useful for detecting non-responders to anti-TNF therapy such as inflixingly therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human;
TNF rec
                                                                                                                                                                                                                                                                             polymorphism II.
                                                                                                                                                                                                                                                                                        Detecting non-responders to anti-human necrosis factor therapy, comprises testing an individual for homozygosity for a single nucleotide polymorphism in the gene coding for the tumor necrosis factor receptor
                                                                       Sequence
                                                                                            the 2 novel polymory inflammatory, maligrepresents a TaqMan
                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                Schreiber S,
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                                                                                                                                                                                                                                                                                                                                                                                       (CONA-)
                                                                                           2 novel polymorphisms are useful for diagnostic purposes in lammatory, malignant or other chronic diseases. The present ir resents a TaqMan probe used in the methods of the present ir
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Pred. No. 2
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0; Mismatches
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Pred. No. 2.2e+03;
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RESULT 3171
ABK69044/c
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XX Alpha 1
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PR 1
                                                                                                                               The invention relates to an isolated beta-related protein (BRP) a novel glycoprotein hormone, its fragment, derivative, analogue, homologue or comaturally occurring allelic variant and the nucleic acid encoding it.

CC Also disclosed are novel alpha related proteins (ARP) and their nucleic caids. Also included are a nucleic acid vector comprising BRP, a host cell comprising the vector, a protein multimer comprising BRP, a host composition, and a second polypeptide, an antibody that selectively binds to BRP or the multimer, screening for a modulator of activity, or of clatency or predisposition to a reproductive disorder comprising CC administering a test compound to an animal at risk from a pathology associated with BRP, where the animal recombinantly expresses an ARP/BRP collective disorder in a subject by measuring the amount of an ARP/BRP polypeptide, measuring the presence of, or predisposition to, a control level, determining the presence of, or predisposition to, a concleic acid in a sample and comparing it to a control and expressing an ARP/BRP protectins, nucleic acids, antibodies and multimers are useful for treating, preventing or diagnosing reproductive and cell proliferative classred including ovulatory diseases, fertility related disorders, they are also useful for stimulating spermatogenesis, increasing the cutous of the thyroid glandular cells, regulating gonadal function, ce function of the thyroid glandular cells, resulating or suppressing certifities, reperfusion or regeneration and treatment of lung or liver fibrosis, reperfusion injury in various tissues and conditions resulting cutous cutous tissues and conditions resulting cutous care disperse and conditions resulting cutous cutous tissues and conditions resulting cutous care in a care and conditions resulting cutous cutous cutous tissues and conditions resulting cutous care in a care and conditions resulting cutous care and condi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel beta subunits useful for treating
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                                          differentiation of tissues from precursor tissues or cells, inhibiting the growth of tissues, for treating inflammatory conditions including
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                                                                                                                    from systemic cytokine damage, for promoting or inhibiting
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       associated with
                                                                             inhibiting
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        This invention relates to the cDNA and protein sequences of novel proteins HYPLIP1 or FCHIL and to sequence variations within these genes that have been shown to be associated with lipid disorders.

Oligonuclectide probes that hybridise to the cDNA sequence are useful for analysing the expression of FCHIL1 by detecting the expression of the mRNA transcript in the sample. A host cell transformed with the cDNA of the invention is useful for producing the protein by recombinant means. Pharmaceutical compositions based on the sequences of the invention are useful for treating or preventing a lipid disorder associated with expression of FCHIL such as familial combined hyperlipidaemia, coronary artery disease, atherogenic lipoprotein phenotype, hyperapobetalipoproteinaemia, hypertriglyceridaemia, familial variational displantation and dyslipidaemia hypertension, syndrome X, obesity, insulin resistance and hyperchalters of the country of the co
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Ohmen J, 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel HYPLIP1 and FCHL1 genes and their sequence variations associated with lipid disorder and cancer, useful for prognosis, diagnosis and
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                                                                                                                                                                                                                                                                                                                                                                                    Claim 11; Page 77; 102pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                          treatment of lipid disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-339808/37.
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The cDNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chatterjee A,
, Wu C;
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Pred.
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No. 2.2e+03;
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the diagnosis
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Best Local S
Matches 16
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08-FEB-2001;
22-FEB-2001;
14-MAR-2001;
                                                                                                                                                                                                                                                                                                                    18-DEC-2000;
21-DEC-2000;
04-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; G protein coupled receptor; GPCR; GPCRX; neuroprotective; nootropic; anti-HIV; antiasthmatic; antiarteriosclerotic; cytostatic; immunomodulator; antiinflammatory; antidiabetic; anorectic; haemostatic; antibacterial; fungicide; protozoal; virucide; nephrotropic; osteopathic; cardiant; antiulcer; antiallergic; hepatotropic; antiparkinsonian; HIV; vaccine; gene therapy; cell signal processing; cardiomyopathy; diabetes; metabolic pathway modulation; atherosclerosis; cancer; obesity; asthma; infection; Parkinson's disease; osteoporosis; Crohn's disease; ulcer; allergy; cirrhosis; glomerulonephritis; stroke; haematopoietic disorder; systemic lupus erythematosus; PCR primer; ss.
New isolated human G-protein coupled receptor x (GPCRX) polypeptide, useful for treating or preventing GPCRX-associated disorders e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          prognosis of predisposition to lipid disorders and cancers, and also identify a molecule which enhances or decreases the HYPLIP1 or FCHLI activity. The present sequence represents an oligonucleotide primer specific for the mouse HYPLIP1 locus of the invention. The mouse HYP locus is situated on chromosome 3
                                                                                                  Smithson
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GPCRX reverse
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                                            WPI; 2002-557660/59.
                                                                                                                                                                                                                         23-MAR-2001;
                                                                                                                                                                                                                                                                                                         10-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2001;
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                                                                                                                                        (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                            12-JAN-2001;
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nes 16; Conser
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                                                                                                            Padigaru M,
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                                                                 adigaru M, Ballinger RA, Kekuda R, Colm
G, Peyman JA, Macdougall JR, Stone D,
E, Millet I, Tchernev VT, Anderson D,
Ellerman KB, Wolenc A;
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; 2001US-0261498P.
; 2001US-0261498P.
; 2001US-0263689P.
; 2001US-0267464P.
; 2001US-0275946P.
; 2001US-0275946P.
; 2001US-0278150P.
; 2001US-0285718P.
; 2001US-0285718P.
; 2001US-029327P.
; 2001US-0312902P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                   2000US-0256635P.
2000US-0257876P.
                                                                                                                                                                                                                                                                                                                                                                            2001WO-US049347.
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Pred.
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No. 2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the invention. The mouse HYPLIP1
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                                                                                                            Colman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
                                                                               .man SD, Sciore P;
Vernet CAM, Shenoy S;
Gusev V, Malyankar UM;
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ABX09454
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cc mootropic, anti-HIV, antiasthmatic, antiarteriosclerotic, cytostatic, cc immunomodulator, antiinflammatory, antidabetic, anorectic, haemostatic, cc antibacterial, fungicide, protozoal, virucide, nephrotropic, osteopathic, cc cardiant, antiulcer, antiallergic, hepatotropic and antiparkinsonian cc activities, and can be used in vaccines and gene therapy. GPCR proteins, cmucleic acid molecules, and antibodies from the present invention can be used for manufacturing a medicament for treating or preventing a GPCR-cc associated disorder or syndrome related to cell signal processing and cmetabolic pathway modulation, such as cardiomyopathy, atherosclerosis, cc diabetes, cancer, obesity, infections (bacterial, fungal, protozoal or viral), HIV, asthma, Parkinson's disease, osteoporosis, Crohn's disease, culcers, allergies, cirrhosis, glomerulonephritis, stroke, systemic lupus erythematosus, or haematopoletic disorders. Anti-GPCR antibodies can be cused diagnostically to monitor protein levels in tissues as part of a clinical testing procedure such as in determining the efficacy of a given treatment regimen. Abg08418 to Abg08639 represent PCR primers and probes cc for the human GPCRs of the present invention
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Best Local
                                                                     Determining genetic risk of arteriosclerosis, for clinical diagnosis, comprises hybridizing patient nucleic acid with an array of probes derived from risk-associated reference genes and their mutations.
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                                           Example
                                                                                                                                   WPI; 2002-723374/78.
                                                                                                                                                                   Cullen
                                                                                                                                                                                                                             13-MAR-2001; 2001DE-01011925
                                                                                                                                                                                                                                                            13-MAR-2002; 2002WO-EP002780
                                                                                                                                                                                                                                                                                                                         WO200272882-A2
                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                   mutation; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                Arteriosclerosis-detecting probe from HCF2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABX09454 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                (OGHA-) OGHAM GMBH
                                                                                                                                                                                                                                                                                            19-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                   Arteriosclerosis;
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Similarity 88.9%;
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                                                                                                                                                                Seedorf U;
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                                                                                                                                                                                                                                                                                                                                                                                                diagnosis; hybridisation; synergism; gene therapy;
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                                      146pp;
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                                           German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8;
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This invention describes a novel method for determining the genetic risk of arteriosclerosis both for clinical diagnosis and for population

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RESULT 3175
ABS97168/c
ID ABS9716
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XX ABS9716
XX Human C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KW cytochrome P450 A2; CYP4501A2; cytochrome P450 CZE; CYP4502E1; LTF;
KW cytochrome P450 A2; CYP4501A2; cytochrome P450 CZE; CYP45002E1; LTF;
KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; MR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW glutathione-S-transferase 1; GST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
KW NADPH quinone oxidoreductase 2; NOO2; sulforransferase thermolabile; STM;
KW NDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7;
KW WT12B7; UDP-glucuronosyl transferase; UGT2B13; urokinase receptor; uPA;
KW multidrug resistance 1; lactotransferain; orphan nuclear receptor;
KW multidrug resistance associated protein 3; cancer; prostate;
KW multidrug muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;
KW acetylcholine muscarinic receptor; cTMMR1; CHMR2; CHMR3; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW central nervous system; pulmonary; immunological; sequencing.
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Best Local S
Matches 16
    Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers
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                                                                                                                             WPI; 2002-698522/75.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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Pred. No. 2.2e+03;
J; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 22;
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В Ś

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disorder-related traits
                           for locating, identifying and characterizing the genes responsible for
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Example 2; Page 101; 714pp; English

CC cyylochrome P450 UZEI (CIP450UZEI), addremented to expect the process of the provide hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (RANT), cathepsin & (CTSS), cyclooxgenase 2 (COX2), diazepam binding CC inhibitor (DBI), epoxide hydrocarbon receptor nuclear translocator grotein (FLAP), glutathione-S-transferase 2 (COX2), bistamine-N-methyl CC (LTAP), glutathione-S-transferase 12 (CST12), histamine-N-methyl CC (LTAP), UDP-Glucuronosyl transferase 2B4 (CC (UGT2B4), UDP-Glucuronosyl transferase 2B4 (CC (MRB1), lactotransferrin (LTF), multidrug resistance associated protein 3 (CC (MRB2), orphan nuclear receptor (NRR112), or acetylcholine muscarinic CC receptor 1, 2, 3, 4, or \$ (CHRR1, CHRR2, CHRR2, CHRR3, CHRR4) crimer e useful as CC (MRB2), orphan muclear receptor locating and characterising the genes that CC genetic linkage markers for locating and characterising the genes that CC drentifying the genes responsible for avariety of disorder-related CC traits as a result of their e.g., overexpression, constitutive CC expression, mutation or underexpression, which may be used in diagnosing CC expression, mutation or underexpression, which may be used in diagnosing the golymorphic sequences contained in CYP4501A1, CYP4501A1, CYP4501A2, CYP4501A2, CYP4501A1, CYP4501A1, CYP4501A1, CYP4501A1, CYP4501A1, CYP4501A1, CYP4501A2, CYP4501A2, CYP4501A2, CYP4501A2, CYP4501A2, CYP4501A3, CYP4501A1, CYP4501A1, CYP4501A2, CYP4501A2, CYP4501A3, CYP4501 human cytochrome P450 Al (CYP4501Al), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1), aryl hydrocarbon (AHR), aryl hydrocarbon receptor betal (ADBR1), the sequence of an isolated nucleic acid

Sequence 22 B₽; 13 A; 3 C; 1 G; 5 T; 0 U; 0 Other;

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                              Query Match
Best Local S
Matches 16
            4459 TGGACTTTTTTTTTTT 4476
18
                               16;
                                      Similarity
                               Conservative
                                      0.2%;
Н
                              ٥.
                                      Score 14.8; DB 1;
Pred. No. 2.2e+03;
                               Mismatches
                                               1;
                                               Length 22;
                                Indels
                                0
                               Gaps
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RESULT 3176

ABK71248 standard; DNA; 22 BP

15-JUL-2002 (first entry

Mouse HYPLIP1 locus PCR primer #321.

Human; mouse; HYPLIP1; FCHL1; familial combined hyperlipidaemia; lipid disorder; PCR; primer; ss.

WO200220848-A2

14-MAR-2002

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RESULT 3177
ABK15346
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                              Mouse; cyclooxygenase-2; COX-2; PCR; primer; sepsis; pancreatitis; l trauma; blood loss; penetrating injury; septic shock; pneumonia; septicaemia; bactaeremia; urinary tract infection; wound infection; septicaemia; bastemic inflammatory response syndrome; PGE_2; prostaglandin E_2; receptor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated polynucleotide comprising a sequence variation of a mouse HYPLIP1 CDNA or a human FCHL1 (familial combined hyperlipidaemia) gene. The FCHL1 polynucleotide, the FCHL1 polypeptide or antibody immunoreactive to the FCHL1 polypeptide are useful for treating or preventing cancer associated with expression of FCHL1, as well as for treating lipid disorder. The mouse HYPLIP1 CDNA or human FCHL1 gene are also useful for diagnosing or prognosing a predisposition to lipid disorder and cancer. ABK70902-ABK71303 represent mouse HYPLIP1, human FCHL1 coding sequences and PCR primers of the invention
Treating a patient at risk for
                                                                                    (STRO/)
                                                                                                                          15-FEB-2000;
                                                                                                                                                  14-FEB-2001; 2001US-00782936.
                                                                                                                                                                           17-JAN-2002
                                                                                                                                                                                                                           an
Mark
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                                                                                                                                                                                                                                                                                                                                                                                                     ABK15346 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 11; Page 77; 102pp; English.
                         WPI; 2002-179019/23.
                                                                                                                                                                                                  US2002006915-A1
                                                                                                                                                                                                                                                                                                                           Cyclooxygenase-2 (COX-2) sense
                                                                                                                                                                                                                                                                                                                                                                              ABK15346;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preventing lipid disorders and cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New mouse HYPLIP1 and human FCHL1 (familial genes and their sequence variations, useful
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                                                Strong VE,
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                                                                                     MACK STRONG
STAPLETON P
                                                                           DALY
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В.
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Wu C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8;
Pred. No. 2.
systemic inflammatory response syndrome
                                                                                                                                                                                                                                                                                                                          PCR primer DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                 Daly JM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8; DB 1;
2.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        combined hyperlipidemia)
for diagnosing, treating
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                                                                                                                                                                                                                                                                                                                               KW human; nuclear factor-kappaB; NF-kappaB; immune disorder; cancer;
KW hiflammatory disorder; apoptosis; hepatic disorder; Hodgkin's lymphoma;
KW haematopoietic tumour; hyper-IgM syndrome; viral infection; asthma;
KW hypohidrotic ectodermal dysplasia; human immunodeficiency virus; HIV;
KW X-linked anhidrotic ectodermal dysplasia; al incontinentia pigmenti;
KW influenza; rheumatoid arthritis; inflammatory bowel disease; colitis;
KW atherosclerosis; cachexia; euthyroid sick syndrome; stroke; EAE;
KW hyperimmental allergic encephalomyelitis; autoimmune disorder; wound;
KW hyper immune activity; acute phase response; hypercongenital condition;
KW hyper immune activity; acute phase response; hypercongenital condition;
KW hyper immune activity; acute phase response; hypercongenital condition;
KW hyper immune activity; acute phase response; hypercongenital condition;
KW wirtam allergic encephalomyelitis; autoimmune disorder; wound;
KW vitamin B12 malabsorption; neurological disorder; diabetes mellitus;
KW vitamin B12 malabsorption; neurological disorder; Huntington's chorea;
KW vitamin B12 malabsorption; virucide; anaemia; antiinflammatory; anti-HIV;
KW vitamin B12 malabsorption; virucide; antiarithratic;
KW vitamin B12 malabsorption; virucide; antiarithratic; cardiant;
KW vitamin B12 malabsorption; virucide; antiarithratic;
KW vitamin B12 malabsorption; virucide; antiarithratic;
KW vitamin B12 malabsorption; virucide; antiarithratic;
KW vitamin B12 malabsorption; viruc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a new method of treating a patient at CC risk for systemic inflammatory response syndrome. The method involves CC administering a selective cyclooxyganase-2 inhibitor or a drug which CC stimulates at least one prostaglandin E 2 (PGE 2) receptor or a drug which interferes with binding of PGE 2 to at least one of PGE 2 complete the invention can be used for treating a patient at risk for cyclooxyganase syndrome e.g. sepsis, pancreatitis, burns, trauma, life threatening blood loss from penetrating injury, or a patient cyclooxyganase syndrome e.g. sepsis, pancreatitis, burns, trauma, life threatening blood loss from penetrating injury, or a patient cyclooxyganase shock, infections such as preumonia, cyclocamia, bactaeremia, urinary tract infection, wound infection or drug reaction and can also be used for beneficial immune modulation. The cyclocamia reduce the incidence of infectious complications and improve trauma, reduce the incidence of infectious complications and improve convival after traumatic injury. The present nucleic acid sequence crepresents the mouse cyclooxygenase-2 (COX-2) sense PCR primer that was used in the invention with the COX-2 antisense PCR primer (ABKI5347) for the isolation and determination of COX-2 mRNA
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                antiarteriosclerotic; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human NF-kappaB associated polynucleotide PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g. trauma involves administering cyclooxygenase-2 inhibitor or a drug.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                 cerebroprotective; cardiant;
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2002WO-US012636

WO200286076-A2

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-JAN-2002; 2002US-0346986P
                                                      10-APR-2003
                                                                                                                                             WO2003029424-A2
                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                immunomodulator; cytostatic; nootropic; neuroprotective; antiparkinsonian; antilipaemic; gene therapy; human disease; metabolic disorder; diabetes; obesity; infection; cachexia; cancer; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human NOVX reverse PCR primer SEQ ID NO:296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; NOVX; antidiabetic; anorectic; antibacterial; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA05936 standard; DNA;
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Pred. No. 2.2e+03;
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2001US-03301042P.
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Smithson G, Millet I, Peyman JA, Kekuda R, Ju J, Li L, Patturajan M, Spytek KA, Edinger SR, Ellerman K, Malyanka Ort T, Gorman L, Zerhusen BD, Anderson DW, Zhong M, Catt Ji W, Miller CE, Rastelli L, Stone DJ, Pena CEA, Shenoy Shimkets RA, Rothenberg ME, Leach MD, Agee ML, Berghs C, Eisen AJ, 'Gangolli EA, Rieger DK, Spaderna SK; Malyankar UM; Catterton E; SG; Dipippo VA;

WPI; 2003-381626/36.

New NOVX polypeptides and nucleic acids, useful for diagnosing, preventing or treating NOVX-associated disorders, e.g. diabetes, obesity, cancer or dyslipidemia, and in chromosome mapping, tissue typing or pharmacogenomics.

Example C; Page 384; 586pp; English.

The present invention describes NOVX proteins, where X can be 1 to 55 CC (e.g. NOV1). Also described: (1) a composition comprising a polypeptide containers, the composition described above; (3) an isolated nucleic acid molecule which encodes a NOVX protein of the invention; (4) a vector CC comprising the nucleic acid molecule described above; (5) a cell comprising the above vector; (6) an antibody that immunospecifically comprising the polypeptide described above; (7) methods for determining the greence or amount of the above polypeptide or nucleic acid molecule in a CC sample; (8) methods for determining the presence of or predisposition to a disease associated with altered levels of expression of the above polypeptide or nucleic acid molecule in a first mammalian subject; (9) a CC method of identifying an agent that binds to the polypeptide described cor cuse in treating a pathology that is related to an aberrant expression or CC use in treating a pathology that is related to an aberrant expression or caberrant physiological interactions of the polypeptide; (11) a method of screening for a modulator of activity or of latency or predisposition to a pathology associated with the polypeptide; (12) a method for modulating the activity of the polypeptide described above; (13) methods of treating or preventing a pathology associated with the above polypeptide in a CC mammal; and (14) a method for producing the above polypeptide. NOVX

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RESULT 3180
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The present invention relates to the isolation of perennial ryegrass (Lollum perenne) nucleic acids and nucleic acid fragments encoding salt inducible or salt responsive proteins. The salt-inducible or salt responsive proteins include salt-inducible proteins (ESI3), glutathione peroxidase homologues (CSA), low-temperature-inducible proteins (LT16), salt-stress induced proteins (SALT), early salt-responding glucose 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   glutathione peroxidase homologue; low-temperature-inducible protein; salt-stress induced protein; SALT; WESRS; ALDP; plant tolerance; salt-stresponding glucose 6 phosphate 1 dehydrogenase; salt shock; plastidic fructose 1,6-biphosphate aldolase homologue; osmotic stress; environmental stress; sodium compartmentalisation; salt stress; pCR; sodium ion influx; sodium ion efflux; plant metabolism; primer; ss.
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                                                                                                                                                                                                                                                                   New substantially purified or isolated nucleic acid or nucleic acid fragment encoding a salt-inducible or salt responsive protein, useffenctic markers for modifying plant tolerance to environmental and/
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RESULT 3181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BODN/)
(CAST/)
(CHAT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse; PCR; primer; ss; HYPLIP1; FCHL1; variation; lipid disorder; allele; anti-lipid disorder; anti-cancer therapy; gene therapy; familial combined hyperlipidaemia; coronary artery disease; atherogenic lipoprotein phenotype; hyperapobetalipoproteinaemia; hypertriglyceridaemia; low density lipoprotein subclass B; LDL; familial dyslipidemic hypertension; syndrome X; hypercholesterolaemia; obsesity; insulin resistance; cancer; cytostatic; antilipaemic; obsesity; insulin resistance; cancer; cytostatic; antilipaemic;
                                                                                                                    Novel isolated polynucleotide comprising a mouse or human combined hyperlipidemia 1 gene having a variation that is a lipid disorder, useful for identifying susceptibility to
                                                                                                                                                                                                                     WPI; 2003-540780/51.
                                                                                                                                                                                                                                                                                          Bodnar
                                                                                                                                                                                                                                                                                                                                                             (ROSS/)
(TAFU/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-JUN-2000;
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LUSIS A J.
OHMEN J.
                                                                                                                                                                                                                                                                                                                                                             ROSS D.
TAFURI S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHATTERJEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CASTELLANI
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oss D, Tafuri
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The invention discloses isolated polynucleotides comprising

mouse HYPLIP1

Claim 11; Page

40; 63pp;

English

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ARESULT 3182
ADB95949
ID ADB9594
AC CYCOST
CYCOST
AC CYCO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic; antilipemic; gene therapy; peptide therapy; HYPLIPI; FCHLI; cancer; metabolic pathway; cellular mechanism; lipid disorder; familial combined hyperlipidaemia; mouse; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                 07-SEP-2001; 2001US-00949427.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-DEC-2003 (first entry)
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                                       (LUSI/)
                                                                                                                                      (BODN/)
(CAST/)
(CHAT/)
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                                                                                                              JONG/)
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                                   LUSIS A J.
OHMEN J.
                                                                                                              JONG
                                                                                                                                          CASTELLANI L
CHATTERJEE A.
                                                                                                                                                                                                              BODNAR J S.
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Pred. No. 2
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RESULT 3183
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                                                                                                                                                                                                                                                                           probe; human papilloma virus; HPV; detection; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 22 BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human FCHLI or mouse screening, peptide therapy
                                      WPI; 2003-432398/41.
                                                                               Lin
                                                                                                                                                                 10-OCT-2001; 2001EP-00123379
                                                                                                                                                                                             16-APR-2003.
                                                                                                                                                                                                                                               Human papillomavirus
                                                                                                                                                                                                                                                                                                      Human papillomavirus type MM8 detection oligonucleotide #9
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                                                                                                          (KING-) KING CAR FOOD IND CO LTD
                                                                                                                                      10-OCT-2001; 2001EP-00123379.
                                                                                                                                                                                                                       EP1302550-A1
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                                                               Lin R,
Shih C,
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                                                                Huang H,
Kao Y,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chatterjee A,
Wu C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1,
Pred. No. 2.2e+03,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                               Lee B,
Pan C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
                                                                  Lee
Chan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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                                                                  P, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 22;
                                                                               Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                             88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Detector for identifying carrier having two parts

human papilloma virus subtypes, comprises carrying first and second oligonucleotides that

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ADC98229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 16
The invention relates to a human G protein coupled receptor (GPCR), BG8 (ADC98222), and nucleic acids encoding it (ADC98221). The human BG8 gene is located on chromosome 12p12. BG8 is involved in controlling the expression of hard keratins such as the type I hair keratins Ha3 and Ha4 The invention also relates to antibodies against BG8, a drug screening method using the antibodies, the compounds identified, and a method for detecting and controlling functional abnormalities in cells producing hard keratin. The invention provides for the treatment and prevention of diseases associated with the function of hard keratin-producing cells, such as hair loss, hair shaft injury, abnormal or insufficient hair growth, diseases of the skin, tongue, nails and hair, and abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention comprises oligonucleotides for detecting and identifying subtypes of human papilioma virus (HPV) contained in a sample. The oligonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an HPV detection oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8kin
                                                                                                                                                                                       Protein for treatment and prevention of diseases associated with the function of hard keratin producing cells, such as hair loss, hair shinjury, abnormal or insufficient hair growth, diseases of the skin, tongue, nails and hair.
                                                                                                                                                                                                                                                                                                                                     03-APR-2002;
20-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                         23-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus
                                                                                                                                                              Example 5;
                                                                                                                                                                                                                                                                                    Inoue
                                                                                                                                                                                                                                                                                                                                                                             02-APR-2003; 2003WO-JP004236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse; murine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC98229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         respectively hybridize with DNA contained in first and second subtypes of the virus.
                                                                                                                                                                                                                                                                                                             (BANY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             se; murine; hard keratin; type I hair keratin; Ha3; BG8; d keratin expression; hair loss; hair shaft injury; hair grown disease; tongue disease; nail disease; hair disease; keratoderma; dermatological; trichological; Taqman analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            musculus
                                                                                                                                                                                                                                                           2003-833733/77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  type
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                                                                                                                                                                                                                                                                                                              BANYU PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
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                                                                                                                                                                                                                                                                                    Nambu
                                                                                                                                                                SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 I hair keratin Ha3 PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 2 A;
                                                                                                                                                                                                                                                                                                                                     2002JP-00100843
2002JP-00369385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IJ
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                                                                                                                                                             ID NO 11;
                                                                                                                                                                                                                                                                                  'n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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                                                                                                                                                                                                                                                                                                              CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                  Shimomura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10
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                                                                                                                                                             67pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 1 T; 0 U;
                                                                                                                                                              Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.
                                                                                                                                                                                                                                                                                  Itadani H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer mHa3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .2e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                    Tanaka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  247R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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hair growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                             Ha3 and Ha4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                 shaft
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ADE47875

ID ADE47875

ID ADE4

XX ADE4

AC ADE4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 16
                                                                                                                                                                               02-AUG-2001;
03-AUG-2001;
08-AUG-2001;
13-AUG-2001;
11-AUG-2001;
11-AUG-2001;
11-AUG-2001;
20-AUG-2001;
21-AUG-2001;
23-AUG-2001;
23-AUG-2001;
23-AUG-2001;
29-AUG-2001;
29-AUG-2001;
29-AUG-2001;
21-SEP-2001;
21-SEP-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2001;
21-AUG-2002;
11-APR-2002;
11-APR-2002;
11-AUG-2002;
11-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      formation of hard keratin such as dot keratoderma. Sequences / ADC98229 represent mouse type I hair keratin Ha3 PCR primers uprobe ADC98227} in Taqman expression analysis of BG8-mediated of type I hair keratin Ha3 in mouse hair follicle tissue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; cardiant; antiarteriosclerotic; hypotensive; immunosuppress. dermatological; anorectic; cytostatic; antidiabetic; haemostatic; anti-HIV; antiasthmatic; antibacterial; virucide; neuroprotective; nootropic; antiparkinsonian; antilipaemic; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE47875;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-SEP-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003076642-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human NOVX forward PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADE47875
                                                                                                                                       (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1892 ACCTCTGCCTCAAGATCA
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16; Conser
                                                                                                                                    CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACCTGTGCCTCCAGATCA
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                                                                                                                                                                               7 2001US - 0311979P

2001US - 03123156

2001US - 0313156

2001US - 0313702P

2001US - 0314702P

2001US - 0314466P

2001US - 0315403P

2001US - 0315803P

2001US - 0315936P

2001US - 0323936P

2001US - 0323936P

2002US - 0354655P

2002US - 0354655P

2002US - 0361764P

2002US - 0373825P

2002US - 0380971P

2002US - 0380971P

2002US - 0380980P

2002US - 0380980P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002WO-US024459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001US-0310951P.
2001US-0311292P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;
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Pred. No. 2.2e+03
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          with
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Zerhusen BD, Pena CEA, S Boldog FL, Vernet CAM, Burgess CE,

1 BD, Patturajan N 1, Shimkets RA, I 7L, Padigaru M, S 7AM, Leite MW, G

Khramtsov

,

Guo

M, Kekuda R, Miller CE, Rieger DK;
Li L, Berghs C, Zhong M, Casman ZJ, Voss
Smithson G, Shenoy SG, Ji W, Gorman L;
Guo X, Anderson DW, Spytek KA, Gerlach VL;
V, Ort T, Ellerman K, Rastelli L, Agee ML;

ΕZ

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                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 3186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cc neuroprotective, nontropic, antiparkinsonian, and antilipaemic activity.

cc A polynucleotide encoding a polypeptide of the invention may have a use

cc in gene therapy, and as a vaccine. A polypeptide of the invention is

cc useful in the manufacture of a medicament for treating a syndrome

cc associated with a human disease, the disease selected from a pathology

cassociated with the polypeptide. These may also be used in diagnosing,

cc treating or preventing NOVX-associated disorders such as cardiomyopathy,

cc therosclerosis, hypertension, scleroderma, obesity, cancer, diabetes,

cc multiple sclerosis, infections, anorexia, cancer-associated cachexia,

cc multiple sclerosis, infections, anorexia, cancer-associated cachexia,

cc multiple sclerosis, infections, anorexia, cancer-associated cachexia,

cc disease), haematopoietic disorders (e.g. Alzheimer's disease or Parkinson's

cc disease), haematopoietic disorders, dyslipidaemias and other wasting

cc disease), haematopoietic disorders, dyslipidaemias and other wasting

cc used as hybridisation probes, in chromosome mapping, tissue typing,

cc used as hybridisation probes, in chromosome mapping, tissue typing,

cc used as yaccines. The present sequence represents a PCR primer used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 16
02-AUG-2001;
03-AUG-2001;
08-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Giot L,
Taupier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chaudhuri A,
Giot L, Ooi
                                                                                                                                                                                                    human; cardiant; antiarteriosclerotic; hypotensive; immunosuppressive; dermatological; anorectic; cytostatic; antidiabetic; haemostatic; anti-HIV; antiaschmatic; antibacterial; virucide; neuroprotective; nootropic; antiparkinsonian; antilipaemic; gene therapy; vaccine; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel (NOVX) human polypeptide. A polypeptide of the invention has cardiant, antiarteriosclerotic, hypotensive, immunosuppressive, dermatological, anorectic, cytostatic, antidiabetic, haemostatic, anti-HIV, antiasthmatic, antibacterial, virucide,
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                                                             02-AUG-2002;
                                                                                                                           WO2003076642-A2
                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                  Human NOVX forward PCR primer SEQ ID NO:240
                                                                                                                                                                                                                                                                                                                  29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                             ADE47878 standard;
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                                                                                                                                                                                         88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptides and nucleic acids, useful for preventing or NOVX-associated disorders, e.g. cancer, diabetes, lerosis, asthma or AIDS, and in chromosome mapping, tissue
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ilarity 88.9%;
Conservative
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2001US-0309501P.
2001US-0310291P.
2001US-0310951P.
                                                               2002WO-US024459.
                                                                                                                                                                                                                                                                                                               (first entry)
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E, Rothenberg
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Pred. No. 2.2e
0; Mismatches
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Spaderna SK, Hjalt T, Liu
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iu X;
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31 AUG-2001;

21-SEP-2001;

03-DEC-2001;

05-FEB-2002;

05-MAR-2002;

19-MAY-2002;

15-MAY-2002;

15-MAY-2002;

28-WAY-2002;

29-WAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-2001;
13-AUG-2001;
14-AUG-2001;
17-AUG-2001;
17-AUG-2001;
20-AUG-2001;
21-AUG-2001;
23-AUG-2001;
28-AUG-2001;
29-AUG-2001;
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                                                                                                                                                                                                                                                                                                      2001US-0311292P
2001US-0311203P
2001US-0313203P
2001US-0313156P
2001US-0313702P
2001US-0314031P
2001US-0315403P
2001US-0315403P
2001US-0315403P
2001US-0315403P
2001US-0315403P
2001US-0315403P
2001US-0315403P
2001US-0315403P
2002US-0361764P
2002US-0361764P
2002US-0361764P
2002US-0380980P
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Boldog FL, P Vernet CAM, Burgess CE, Chaudhuri A, Giot L, Ooi Giot L, Oc Taupier RJ, Zerhusen BD, Patturajan M, Kekuda R, Miller CE, Rieger DK;
, Shimkets RA, Li L, Berghs C, Zhong M, Camman SJ, Voss I
L, Padigaru M, Smithson G, Benoy SG, Ji W, Gorman Li
LAM, Leite MW, Guo X, Anderson DW, Spytek KA, Gerlach VL;
CE, Khramtsov NV, Ort T, Ellerman K, Rastelli L, Agee ML;
i A, Chant JS, Dipippo VA, Edinger SR, Eisen A, Gangolli I
Ooi CE, Rothenberg ME, Spaderna SK, Hjalt T, Liu X; CE, Rothe M EΑ EZ;

WPI; 2003-779062/73

New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, diabetes, atherosclerosis, asthma or AIDS, and in chromosome mapping, tissue typing pharmacogenomics.

Example SEQ ID NO 240; 562pp; English

The invention relates to a novel (NOVX) human polypeptide. A polypeptide of the invention has cardiant, antiarteriosclerotic, hypotensive, communosuppressive, dermatological, ancrettic, cytostatic, antidiabetic, antimunosuppressive, anti-HIV, antiasthmatic, antibacterial, virucide, anti-HIV, antiasthmatic, antibacterial, virucide, communosuppressive, nootropic, antiparkinsonian, and antilipaemic activity. Comeuroprotective, nootropic, antiparkinsonian, and antilipaemic activity. Compared the conding a polypeptide of the invention may have a use in gene therapy, and as a vaccine. A polypeptide of the invention is cuseful in the manufacture of a medicament for treating a syndrome community of the community of the polypeptide. These may also be used in diagnosing, compassing or preventing NoVX-associated disorders such as cardiomyopathy, cascociated with the polypeptide. These may also be used in diagnosing, community of the polypeptide of the invention of the invention of the polypeptide of the invention of the invention and the cachexia, cascociated with the polypeptide. These may also be used in diagnosing, community of the invention of the associated cachexia, concerned the manufacture of the invention of the useful as vaccines. the invention.

Sequence 22 ₽₽; 1 A; 11 Ç 0 ູດ 10 ; 0 U; 0 Other;

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                 Matches
                         Query Match
Best Local
                16;
                         Similarity
                 Conservative
                         0.2%;
                0;
                         Score 14.8; DB 1; Length 22; Pred. No. 2.2e+03;
                        .2e+03
                  Indels
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                 Gaps
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RESULT 3188
AAN70275/c
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AAN70276/c
                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                         The patent claims a new molecule of formula (NA1---S---NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; --S--- = a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolyable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        )276/c
AAN70276 standard; DNA; 26
                                                                           03-OCT-2002
26-MAY-1991
Synthetic
                       Hybridisation; probe; ss.
                                                  Sequence
                                                                                                                                           AAN70275
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26-MAY-1991
                                                                                                                   AAN70275;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-DEC-1985;
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                                                                                                                                                                                                                                    4012
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                                                                                                                                           standard;
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fi
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                                                                                                                                                                                                                                                             0.2%;
llarity 73.1%;
Conservative
                                              scissile link probe MRC059
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                                                                          (revised)
(first entry)
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(first entry)
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                                                                                                                                                                                                                                                                                                                 A; 0 C; 0
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                                                                                                                                                                                                                                                            Score 14.8; DB 1;
Pred. No. 2.6e+03;
0; Mismatches 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Robertson
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                                                                                                                                                                                                                                                                                                                 0 Other;
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RESULT 3189
AAN92241/c
ID AAN92241 standard;
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Best Local Similarity
                                                                                                                                                                                                                                                                                      25-MAR-2003
31-OCT-2002
25-APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The patent claims a new molecule of formula (NA1---S---NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S--- = a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic linked by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP227976-A.
           02-NOV-1989
                                     WO8910415-A.
                                                                                      misc_feature
                                                                                                                           misc_feature
                                                                                                                                                               misc_feature
                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                   Probe MRCO59; solid
                                                                                                                                                                                                                                                             SS
                                                                                                                                                                                                                                                                                                                                           AAN92241;
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                                                                                                                                                                                                                                                            probe MRCO59.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid probes a scissile linkage.
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(first en
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                                                                                     /note= "ribonucleotides."
|5. .26
                                                                                                            /*tag=
                                                                                                                             /note= "deoxyribonucleotides."
11. .14
                                                                                                                                                                             Location/Qualifiers
                                                            note= "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                      DNA; 26
                                                                                                                                                                                                                                   support; ribonuclease.
                                                                                                                                                                                                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                                                                       BP.
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Pred. No. 2.6e
0; Mismatches
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RESULT 3190
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Best Local Similarity
Matches 19; Conser
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31-OCT-2002
25-APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         probe MRCO59 is bound by a hydrolysable linkage to a solid support at its 3' end. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least once within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. Using this system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the ribonucleotides by a ds RNase, eg RNase H or Excil. (Updated on 31-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                                                                                                                                                                          misc_feature
                                                                                                                                                                                                                                                                                                              SS probe MRCO60.
                                                                                                                                                                                                                                                                                                                                                                                           AAN92242;
                                                                                                                                                                                                                                                                                                                                                                                                                   AAN92242 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 26 BP; 0 A; 0 C; 0 G; 22 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 24; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1989-339977/46
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            29-APR-1988;
                                     29-APR-1988;
                                                                                                                                        misc_feature
                                                                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                            Synthetic
                                                              02-NOV-1989
                                                                                                                                                                                                                                                                                    Probe MRCO60; solid support; ribonuclease.
                                                                                        WO8910415-A
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                                                                                                                                                                                /note= "deoxyribonucleotides.
13. .16
                                                                                                                                          /note= "ribonucleotides.
17. .26
                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                   /*tag=
                                                                                                                 note= "deoxyribonucleotides
                                                                                                                                                                                                         *tag=
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Pred. No. 2
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RESULT 3191
AAF77536/c
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3' end. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least once within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. Using this system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the ribonucleotides by a ds RNase, eg RNase H or ExoIII. (Updated on 31-OCT-2002 to add missing OS field.)
                                                                                                                                                                (LINS/) LIN S. (CHUO/) CHUONG (YING/) YING S.
                                                                                                                                                                                                                                                                                                                                                           cDNA library production; SCLA; gene chip technology; differential screening; pathological diagnosis; genetic identification; single-cell cDNA library amplification; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Probe MRCO60 is bound by a hydrolysable linkage to a solid support at
3' end. It is used by reacting excess probe with a target nucleic action
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26 BP; 0 A; 0 C; 0 G; 22 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 24; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Duck P,
                                                 Generating a complete full-length cDNA library from single cells for use in gene chip technology, involves reverse transcribing intracellular mRNAs, adding polynucleotide tail and amplifying formed cDNAs.
                                                                                                                                                                                                                                                                             06-MAR-2001.
                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA library
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                                                                                                            WPI; 2001-243448/25.
                                                                                                                                      Lin S,
                                                                                                                                                                                                                                                  20-NOV-1998;
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Pred. No. 2
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The present invention describes a method of producing full-length cDNA

Disclosure; Col 11-12; 11pp; English

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RESULT 3192
AAD03682/c
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Best Local Similarity
Matches 19; Conserv
The patent discloses novel human phosphodiesterase (PDE), zcytor13 cDNA and its corresponding protein. Zcytor13 protein is used to promote wound healing in tissues, to exhibit anti-bacterial and anti-viral effects and to identify modulators (e.g. agonists or antagonists). Zcytor13, its agonists or antagonists or disclamation, inflammation, during and after surgery, arthritis, asthma, inflammation, bowel disease or diverticulitis, for modulating spermatogenesis, sperm capacitation, as immunocontraceptive or anti-fertility vaccine and for treating male infertility. Zcytor13 protein and its antibodies are used to diagnose cancer, reperfusion ischaemia, asthma, psoriasis and melanoma. Zcytor13 proteins are used to enhance fertilisation. Zcytor13 antagonists are used to treat myocarditis, atherosclerosis, pelvic inflammatory diseases. Zcytor13 psoriasis, eczema, scleroderma and other inflammatory diseases. Zcytor13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antipsoriatic; cytostatic; antiatherosclerotic; antiinfertility; cardiant; antiinflammatory; dermatological; wound healing; antiviral; antibacterial; therapy; inflammatory bowel disease; diverticulitis; spermatogenesis; sperm capacitation; immunocontraceptive; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          libraries from single cells, designated single-cell cDNA library amplification (SCLA). The method is useful in gene chip technology, differential screening, pathological diagnosis, physiological prognosis and genetic identification. No further information about this sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoding it, modulators a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer; reperfusion ischaemia; psoriasis; melanoma; myocarditis; PID; pelvic inflammatory disease; eczema; scleroderma; vasoconstriction; heart arrhythmia; congestive heart disease; muscle spasm; fatigue; chromosomal abnormality; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1C; Page 118; 122pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-OCT-2000; 2000WO-US027734.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human phosphodiesterase polypeptide, zcytor13 and polynucleotide ing it, for detecting human chromosomal abnormalities, identifying ators and treating inflammatory and cardiovascular diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         full length
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Pred. No. 2
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AAF23526/c
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Matches 19
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           Query Match
Best Local
                                                                                                                                                                                                                                                                            (LINS/) LIN S.
(YING/) YING S.
(CHUO/) CHUONG C.
(WIDE/) WIDELITZ I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequences and/or its antibodies are useful for treatment of disorders associated with vasoconstriction, heart arrhythmia, congestive heart disease, muscle spasms and fatigue. They are used for detecting human chromosomal abnormalities. Zcytor13 cDNAs are used in gene therapy.
                                                Sequence
                                                                      transcription, denaturation, doutranscription. The invention is limited mRNAs from single cells
                                                                                                                                                                         Generating amplified messenger RNA sequences from single cells, involves cycling steps of reverse transcription, denaturation, double-stranded DNA sequences and in vitro transcription.
                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                     Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primer; mRNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence is a polyA PCR primer, ZC7764b which is used to isola length zcytor13 cDNA by screening human placental cDNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zcytor13-cytokine fusion proteins or antibody-cytokine fusion proteins are useful for enhancing in vivo killing of target tissue. The present sequence is a polyA PCR primer, ZC7764b which is used to isolate full
                                                                                                            polymerase reaction activity,
                                                                                                                                                                                                                                                                                                                                                                    04-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                             14-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                     WO200075356-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF23526
                                                                                     present invention relates to generating amplified messenger RNAs with present invention relates to generating amplified messenger RNAs with ymerase reaction activity, comprising cycling steps of reverse necription, denaturation, double-stranded cDNA synthesis and in vitro nscription. The invention is used for generating amplified mRNAs from
                                                                                                                                                                                                                                                     ß
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             Similarity
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                                                 BP; 0
                                                                                                                                                  Page 17;
 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplification; 88
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73.1%;
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Score 14.8; I
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                       DB 1;
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RESULT 3195
ABS52638/c
ID ABS52638 standard; DN
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AC ABS52638;
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DT 15-NOV-2002 (first of the text)
CT 15-NOV-2002 (first of text)
CT XX
CE Human secreted salivative
XX
CE Human; secreted salivative
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CE Human; secreted salivative
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AAS20596/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a polynucleotide derived from the 4q12-4q13 region of human chromosome 4 and encoding a zsig63 polypeptide, a secreted salivary protein with anti-microbial activity. Due to their microbial activity, the sequences can be used in the study of microbial infections, e.g. for recombinant production of anti-microbial proteins. The sequences can be used in the treatment of tooth decay, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, skin infections, epithelial wounds, chronic tissue damage, acquired immunodeficiency syndrome (AIDS), cystic fibrosis, lung infections, sarcoidosis, emphysema and chronic bronchitis. This sequence represents a sequencing primer for cDNA encoding human zsig63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; zsig63; chromosome 4q12-4q13; salivary protein; antimicrobial; microbial infection; tooth decay; periodontal disease; thrush; emphyses gastrointestinal disease; urinary tract infection; vaginal infection; skin infection; epithelial wound; chronic tissue damage; cystic fibros: acquired immunodeficiency syndrome; AIDS; lung infection; sarcoidosis; chronic bronchitis; gene therapy; protein therapy; primer; ZC7764a.
     Human; secreted salivary protein; zsig63; immunogen; zsig63-cytokine;
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                                      Human secreted salivary protein zsig63 PCR primer ZC7764a.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polynucleotides encoding salivary proteins useful as anti-microbial
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                                                                                                                                                                                                                                                                                                                                                                              Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U;
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                                                                           (first entry)
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                                                                                                                                                DNA;
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The present invention relates to a new secreted salivary protein, zsig63. CC The invention is useful for detecting in a test sample, the presence of calso useful as an immunogen for producing an antibody to zsig63 cc also useful as an immunogen for producing an antibody to zsig63 cc polypeptide. zsig63-cytokine fusion proteins or antibody-cytokine fusion protein are useful for enhancing in vivo killing of target tissues. Cc Pharmaceutical composition comprising purified zsig63 polypeptide are useful in the treatment of conditions associated with pathological complexitions associated with pathological complexitions and viral infections. High cc expression of zsig63 in salivary gland suggests that anti-microbial periodontal disease, thrush and gastrointestinal disease. Other capalications can be used in urinary tract infections, vaginal infections, periodontal disease, thrush and gastrointestinal disease. Other capalications can be used in urinary tract infections, vaginal infections, periodontal disease colonisation, and invasion. The invention is useful when pro-inflammatory activity is desired. Applications for such pro-inflammatory activity is desired. Applications for such pro-capacitic particularly in areas having a limited or damaged vascular system e.g., and the properties may be useful as anti-inflammatory agents. The invention is useful for the treatment of patients having incompetent immune system, such as AIDS (acquired immunodeficiency syndrome) patients or individuals that have undergone chemotherapy, radiation treatment. The invention is useful first material for the treatment of lung infections associated with cystic fibrosis and its agonists or antagonists are useful for aiding digestion. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention for identification of zsic63
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel secreted salivary protein, zsig63 and polynucleotide encoding it useful for treating microbial infections, inflammatory conditions, dental caries and lung infections associated with cystic fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 29; 33pp; English.
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4011 TAAAATGAGAAAAAAGAGAGAAAACA 4036
                                                                                                                                                                                                                                                                                                     methods of the invention for
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SHEPPARD P O.
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                                                                                                                                                                                                                                                                                                     identification of zsig63
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as zsig63 and nucleic acid molecules encoding such polypoptides. Zsig63 can be used in detecting agonists and antagonists of its activity, and is also useful as a host defense polypoptide, immune modulating factor, cc also useful as a host defense polypoptide, immune modulating factor, cc cytokine, or as secreted extracellular matrix associated proteins with cc growth factor hormone activity. It is useful for treating conditions cc associated with pathological microbes, including bacterial, fungal and cviral infections, for treating dental carries (tooth decay), periodontal clients, vaginal infection and for preventing infection in skin and cother epithelial wounds zsig63 is useful for establishing normal context grainst pathogenic colonisation and invasion, for treating chronic tissue damage e.g. damage in extremitles associated with disease, for treating dependence of lung dysfunction, salivary gland dysfunction, or dysfunction of corresponded associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human traits and as considered with disease or other human from the disported and human zsig63 conhibiting chondrosarcomas, atherosclerosis, restenosis and obesity. Th
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AAD45055/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel secreted salivary polypeptide, zsig63, useful as antimicrobial agent for treating microbial infection, dental carries, periodontal disease, thrush gastrointestinal disease, and for aiding digestion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune modulating factor; antipathogenic; cell-cell signalling molecule growth factor; cytokine; growth factor hormone activity; dental carries infection; tooth decay; periodontal disease; growth factor infection; diabetes; obesity; anti-inflammatory; chronic tissue damage; lung dysfunction; restenosis; gene therapy; salivary gland dysfunction; prostate gland dysfunction; forensic DNA profiling; chondrosarcoma; atherosclerosis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to human secreted salivary polypeptide designated as zsig63 and nucleic acid molecules encoding such polypeptides. zsig63
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17-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       secreted salivary protein;
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2000US-00527345.
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Sequence

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Query Match Best Local S Matches 19

Similarity

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Score 14.8; Pred. No. 2

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, Matches' 19
The present invention relates to the isolation of a novel cytokine, CC zalphall Ligand and the polynucleotide encoding it. The invention also CC gives the sequence for the zalphall receptor and the polynucleotide CC encoding it. The zalphall Ligand polypeptide stimulates proliferation of CC entoding it. The zalphall Ligand polypeptide stimulates proliferation of NK CC cells, proliferation of T-cells or NK cell progenitors, the activation of NK CC cells, proliferation of T-cells, proliferation of B-cells stimulated with anti-IgM antibodies. The CC anti-CD40 antibodies, stimulates an antigenic response in a mammal, and CC reduces proliferation of B-cells stimulated with anti-IgM antibodies. The CC zalphall Ligand polypeptide is also useful in preparing antibodies that bind to zalphall Ligand epitopes. The zalphall Ligand polypucleotides can be used as probes or primers to clone regions of a zalphall Ligand gene, and in gene therapy. Zalphall Ligand may also be used to identify CC inhibitors of its activity to enhance the generation of anti-tumour responses with or without the infusion of donor lymphocytes, and to activate or stimulate the immune system. The present sequence represents
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11-MAR-1999;
01-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                             New zalphall ligand polypeptides and polynucleotides, useful for
stimulating proliferation, activation, differentiation and/or of
of inhibition of specialized cell function, or for stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novak JE,
Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytokine; zalphall Ligand; zalphall receptor; NK cell prognatural killer cell proliferation; T-cell proliferation; B-cell proliferation; anti-tumour response; immune system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-MAR-2000; 2000US-00522217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human zalphall Ligand sequencing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ZYMO ) ZYMOGENETICS INC
                                 sequencing primer used to 
uman zalphall Ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4011 TAAAATGAGAAAAAAAAGAGAGAAAAACA 4036
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                                                                                                                                                                                                                                                                                                                                             Col 139; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Presnell SR,
Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0123547P.
99US-0123904P.
99US-0142013P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytostatic; human; sequencing primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sprecher CA, Foster DC,
Nelson AJ, Dillon SR,
                                                sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8; D
Pred. No. 2.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer ZC7764a.
                                                 CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.6e+03;
                                                   clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hammond
                                                 in the isolation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell progenitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                   induction
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ABZ24784/c
ID ABZ247
XX
AC ABZ247
                                                                                                                                                                                                                 RESULT 3198
AAD43853/c
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                                                        RESULT 3199
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                                                                                                                          á
                                                                                                                                                          Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                       The invention relates to a method for obtaining a single stranded polynucleotide tag from a biological sample by cleaving one of the complementary strands of a double stranded polynucleotide with a cleavage agent capable of recognising a double stranded polynucleotide comprising complementary strands and cleaving only one of the strands of the polynucleotide in the process of generating a single stranded polynucleotide tag. The method is useful for separating, analysing tags originating partly, and preferably wholly from a source of DNA and/or RNA in a sample comprising biological cells. The method is particularly for analysing gene expression (expression profiling or differential gene expression), or in diagnosing clinical conditions. The present sequence is a primer used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Obtaining single stranded polynucleotide tags from a biological sample, for analyzing gene expression or diagnosing clinical conditions, comprises employing nicking endonucleases that cleave complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JAN-2001; 2001DK-00000126.
12-FEB-2001; 2001US-0267704P.
 ABZ24784;
                             ABZ24784 standard; DNA; 26 BP
                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-636542/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-JAN-2002; 2002WO-DK000052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer #2 used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-NOV-2002
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                                                                                                                          4012 AAAATGAGAAAAAAGAGAGAAAAACAA 4037
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 294; 302pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAAAATGAGAAAAAAGAGAGAGAAAACA 4036
                                                                                                    AAAAAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                        0.2%;
nilarity 73.1%;
Conservative
                                                                                                                                                                                                                 BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EXPRESSION APS
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                                                                                                                                                                                                                 0 C; 0
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                                                                                                                                                           <u>,</u>
                                                                                                                                                                                                                   <u>ن</u>
                                                                                                                                                        Score 14.8; D
Pred. No. 2.6e
0; Mismatches
                                                                                                                                                                                                                   26 T; 0 U;
                                                                                                                                                                         2.6e+03;
                                                                                                                                                                                   DB 1; Length 26;
                                                                                                                                                                                                                   0 Other;
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                                                                                                                                                           Gaps
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TXXXXXXX
                                                                                                     RESULT 3200
ABX93599/c
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                                                                                                                                                                                                              Matches
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Best Local :
                                                                                                                                                                                                                                                                                                      The present sequence is that of a thiosubstituted oligodeoxynucleic acid (ODN) molecule, ODN 24, including deoxynutidine monophosphates. The invention is based on the discovery that ODNs containing deoxynutidine residues (U-ODNs) have an immunostimulatory effect comparable to, or in many instances greater than, ODNs containing to motifs, producing higher numbers of specific T cells to given antigen. The U-ODNs do not induce the systemic production of pro-inflammatory cytokines and, in contrast to CPG ODNs, are not dependent on a specific motif or a palindromic sequence. Use of a U-ODN for the preparation of a vaccine is claimed. Combining the U-ODN with an antigen strongly increases the potential of the antigen to raise the protection/immune response of a vaccinated individual. An example of the invention demonstrated the generation of a specific immune response against a melanoma-derived peptide (see
                                                                                                                                                                                                                                                                                             ABP58360)
24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligodeoxynucleic acid molecule ODN
                                                                                                                                                                                                                                                                 Sequence 26 BP; 0 A; 0 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-183880/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INTE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAY-2001; 2001AT-00000805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-MAY-2002; 2002WO-EP005448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-APR-2003
           Human zsig63 PCR/sequencing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 8; Page 32; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New oligodeoxynucleic acid molecules useful for the preparation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lingnau K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200295027-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulant; oligodeoxynucleic acid; ODN; vaccine; DNA-RNA hybrid;
                                     28-MAY-2003
                                                                                          ABX93599 standard; DNA; 26 BP
                                                                                                                                                                                     4012 AAAATGAGAAAAAAGAGAGAAAACAA 4037
                                                                                                                                                                                                            al Similarity
19; Conserv
                                                                                                                                                            26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INTERCELL BIOMEDIZINISCHE FORSCHUNGS CISTEM BIOTECHNOLOGIES GMBH.
                                                                                                                                                            Schellack C,
                                                                                                                                                                                                               Conservative
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                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "thiophosphate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag=
                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schmidt W;
           primer 2C7764a.
                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                   G; 1 T; 25 U; 0 Other;
                                                                                                                                                                                                                            Score 14.8;
Pred. No. 2
                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24.
                                                                                                                                                                                                                             6e+03
                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                        Length 26;
                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                 Gaps
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lung infection; cystic fibrosis; lung dysfunction; digestive; salivary gland carcinoma; becamecystis carinii infection; emphysema; chronic bronchitis; prostate dysfunction; prostate adenocarcinoma; cell culture media; gene therapy; human chromosome 4q12-4q13;
                                                                                                                                                                                                                                                                                              ss; PCR; zsig63; adhesin; salivary gland; dental carries; periodontal disease; thrush; gastrointestinal disease; epithelial wound; urinary tract infection; vaginal infection; skin infection; primer; pro-inflammatory; chronic tissue damage; vascular system; diabetes; AIDS;
dentinogenesis imperfecta; dentin dysplasia type II.
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US2002173027-A1

21-NOV-2002

03-AUG-2001; 2001US-00922469

17-MAR-1999; 17-MAR-2000; 99US-0124820P. 2000US-00527345.

(ADLE/) SHEPPARD P O ADLER D A.

Adler DA, Sheppard PO

WPI; 2003-328428/31.

Novel isolated zsig63 polypeptide, member of the adhesin family, useful for treating dental carries, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, infections.

1; Page 29; 32pp; English

cc vector, a cultured cell comprising the vector and expressing the protein, cc a DNA encoding a fusion protein (comprising amino acids 1-15, 16-37, 38-126, 127-219 or 16-219 of zsig63 and an additional protein), using a CC zsig63 reporter gene construct to identify zsig63 agonists, and producing cc an anti-zsig63 antibody using zsig63 immunogenic peptides, Zsig63 is CC useful for detecting in a test sample, the presence of antagonist of zsig63 protein activity. Zsig63 has antimicrobial activity and since cexhibits high expression in salivary gland, can be used for treating CC dental carries, periodontal disease, thrush, and gastrointestinal CC dental carries, periodontal disease, hurush, and gastrointestinal CC disease, urinary tract infections, vaginal infections, skin infections can deter epithelial wounds. The polypeptides can be used for treating CC invasion. Zsig63 can also be used for providing pro-inflammatory activity correct invasion. Zsig63 can also be used for providing pro-inflammatory activity correct invasion. Zsig63 can also be used for providing pro-inflammatory activity correct invasion. Zsig63 patients or in individuals that have undergone consumterapy, radiation treatment, for treating lump infections e.g. in cystic fibrosis. Detection of zsig63 polypeptide at relatively high conditions associated with salivary gland or lung dysfunction including conditions associated with salivary gland or lung dysfunction including conditions associated with salivary gland or lung dysfunction including conditions and may be used to replace serum that is commonly used in culture. The DNA is unseful in gene therapy applications to increase or inhibit zsig63 activity, and for detecting abnormalities on human conditions and dentin dysplasia type III. Zsig63 is an addesin family member. The correct in the conditions and for detecting is an and serious conditions. The invention relates to an isolated zsig63 polypeptide comprising at least 90% identity to an amino acid sequence which comprises domain 1 zsig63, domain 2, domain 3, mature zsig63 and full length zsig3. Also included are the polynucleotide encoding zsig63, a zsig63 expression present sequence is a primer used encoding human zsig63 to isolate and sequence nucleic acdis Of.

Sequence 26 BP; 1 A. 0 Ç 0 ଦ 25 H. 0 U; 0 Other

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                                                                                                                                                                                                                                                                                                                                                 RESULT 3201
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                       ACA62282 standard; DNA; 26
                                                                                                                                                                                                           Lin S,
                                                                                                                                                                                                                                 25-JAN-2000; 2000US-00494212
                                                                                                                                                                                                                                           07-SEP-2001; 2001US-00949305.
                                                                                                                                                                                                                                                                 US2003022318-A1.
                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                      ss; PCR; primer; ant promoter containing
                                                                                                                                                                                                                                                                                                      Oligo (dT) primer #1.
                                                                                                                                                                                                                                                                                                                            ACA62282;
                                                                                                                                                                                                WPI; 2003-479488/45
                                                                                                                                                                                                                      (EPIC-)
                                                                                                                                                                                                                                                                                                                 12-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                            4011 TAAAATGAGAAAAAAGAGAGAAAACA 4036
                                                                                                                                                                                                                                                                                                                                                                  26
                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                      EPICLONE INC
                                                                                                                                                                                                                                                                                                                                                                  TARAAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                      antisense therapy; mRNA expression profile; ing primer.
                                                                                                                                                                                                                                                                                                                                                                                             73.1%;
                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; D
Pred. No. 2.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                             2.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                  Length 26;
                                                                                                                                                                                                                                                                                                                                                                                        Indels
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Improved polymerase thermocycling reaction for nucleic acid amplification, by thermal cycling of promoter-linked nucleic acid template synthesis and in vitro transcriptional amplification of nucleic acid sequences.

Example 4; Page 14; 28pp; English

CC (MI) for linear amplification of nucleic acid sequences, involves and denatured (I) with a promoter acid templates (I), combining the CC denatured (I) with a promoter acid templates (I), a primer (P2), a CC number of deoxymucleotide triphosphates and ribonucleotide triphosphates, CC number of deoxymucleotide triphosphates are reverse transcription enzyme, a DNA-dependent DNA polymerase and RNA CC polymerase, contacting P1 with (I) to generate a number of promoter-containing templates, contacting P2 with the denatured promoter-containing templates to CC generate a number of promoter-containing templates to CC generate a number of promoter-containing double-stranded DNA templates are flanked by P1 in one CC end and P2 in the other end of the other orientation, transcribing the promoter-containing double-stranded DNA templates to form a number of CC promoter-containing double-stranded DNA templates to form a number of containing double-stranded DNA templates to form a number of containing double-stranded DNA templates to form a number of promoter-containing double-stranded DNA templates, contacting the amplified RNA containing double-stranded DNA templates, to form a number of containing the amplified RNA complified RNA templates, contacting the amplified RNA complified RNA templates, contacting the machine of in uncleic acid sequences, and thus for producing mRNA complified RNA is also useful for determining aberrant protein production of cells in a diseased state, by generating an expression profile by the capression profile of the cells in the normal and diseased states, comparing the contacting the mRNA composition of the cell(s) in the diseased state, by the diseased state, amplifying the mRNA sequences of cell(s) in the diseased cate that differ from mRNA in cell(s) in non-diseased state, amplifying the solated state, amplifying the solated state, amplifying aberrant protein function of the cell state and diseased states. The invention relates to an improved polymerase thermocycling reaction amplifying

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AAV71935/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell in a diseased state caused by aberrant protein production, by

determining protein expression of a cell in a diseased state, determining

the mRNA sequence for the aberrant proteins, synthesising an antisense

sequence of the mRNA, amplifying the antisense mRNA sequences by MI, and

comprising the anti-sense mRNA and a compatible lipid based biological

comprising the anti-sense mRNA and a compatible lipid based biological

comprising the aberrant protein, by determining aberrant protein

production of cell in a diseased state by the above method, amplifying

the aberrant protein by MI and using recombinant techniques to determine

the effect of proposed drug on the aberrant protein. MI is also useful

for differential screening of tissue-specific gene expression at a

cellular level, for preparing labeled RNA/DNA probes for a gene chip

technology, and for determining the efficacy of a drug regiment against a

gene or its cDNAs. The present sequence is an Oligo (dT) primer used to

produce second strand cDNA in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                              13-MAY-1997;
19-MAY-1997;
27-MAR-1998;
The invention relates to preparation of a normalised, subdivided library of amplified cDNA from the coding regions of mRNA in a sample. The method involves reverse transcription, with at least one cDNA primer of formula 5'-Con1-dTn2-Vn3-Nn4 to form first stand cDNA where Con1 = any sequence of 1-100 nucleotides; dT = deoxythymidiny1; n2 is at least 1; n3 and n4 are both 0, or n3 is 1 and n4 is at least 1; followed by second strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Normalised; cDNA library; mRNA cloning; reverse transcription; immobilise; screening; hybridisation; nucleic acid amplification; expression pattern; drug development; PCR primer; RT-PCR; ss.
                                                                                                                           Example
                                                                                                                                                                               Preparation of normalised, subdivided cDNA libraries from mRNA - reverse transcription and amplification, used to screen for new of
                                                                                                                                                                                                                                                                                                                                                                                                                   13-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-NOV-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9851789-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anchored poly T RT-PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV71935 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein coded for by the isolated mRNA. M1 is also useful for treating a
                                                                                                                                                                                                                                     WPI; 1999-009772/01.
                                                                                                                                                                                                                                                                                                          (DISP-) DISPLAY
                                                                                                                                                            interacting proteins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4012 AAAATGAGAAAAAAGAGAGAAAACAA 4037
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                                                                                                                           1; Page
                                                                                                                                                                                                                                                                        PR
                                                                                                                                                                           transcription and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                            97DK-00000547.
97US-00871030.
98DK-00000432.
                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-DK000186
                                                                                                                         29; 71pp;
                                                                                                                                                                                                                                                                                                            SYSTEMS BIOTECH
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                                                                                                                                                            potential drugs,
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                                                                                                                           English.
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Pred. No. 2.
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                                                                                                                                                              and for diagnosis.
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RESULT 3203
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA synthesis using the first strand as template and a second cDNA primer of a similar formula, in the presence of DNA polymerase I (or its Klenow fragment) and amplification of double-stranded cDNA with a set of amplification primers. Comparison of cDNA in the prepared library with a database (a computer-generated list of molecular weights of restricted DNA fragments of known sequence) is used to determine presence of an expressed protein in a cell, also to detect changes in such expression (particularly for diagnosis of disease). Surfaces (chip) having amplified cDNA stably immobilised on it, obtained by a similar method, are used to screen for genes of a particular family, by hybridisation with nucleic acid from the family (to identify new genes) and to detect differences in expression patterns between cells. The polypeptides expressed by the libraries can be used for drug development. Sequences AAV/1935 to
                                                                                                                                                                                                                                                                             Inserting a nucleic acid into a circular vector comprising joining their ends, melting, and reannealing ends at two different concentrations, useful for cloning small amounts of nucleic acids and forming genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-442381/38.
                                                                                                                                                                                                                                                                                                                                                                                        Romantchikov Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                         (ROMA/) ROMANTCHIKOV Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4011 TAAAATGAGAAAAAAAGAGAGAAAAACA 4036
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Pred.
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This invention describes a novel method (M1) for inserting a nucleic acid (N1) into a circular vector (V1) comprising joining ends of N1 and V1 under a first nucleic acid concentration, melting hybridized cohesive circularization ends, and reamnealing the ends at a second concentration. The methods are useful for the cloning small amounts of nucleic acids and forming genomic libraries of complex oppulations of DNA or cDNA. The methods allow the cloning of minute amounts of nucleic acids efficiently and avoids the size selection problems of prior art systems. Larger representative libraries to be made. Vector to vector ligation is avoided using the methods. AAA40351-A40366 represents primers used to illustrate the method of the invention

is avoided

Example 3;

Page 67; 71pp; English.

acid

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RESULT 3204
AAA57856/c
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catalytically inactive, but which becomes catalytically active in the presence of an analyte (the effector); providing a nucleic acid substrate for the catalytic activity of the nucleic acid sequence; and contacting the nucleic acid sequence and the substrate with the sample under conditions allowing catalytic activity of nucleic acid sequences. The catalytic nucleic acid sequence will be able to convert the nucleic acid substrate into a nucleic acid product only if the analyte of interest is present. The nucleic acid product only if the analyte of interest is gresent. The nucleic acid catalytic product is then amplified, and a significant increase in the amount of product indicates the presence of the analyte determination of an analyte in a sample in diagnostic quantitative determination of an analyte in a sample in diagnostic assays. The invention describes the in vitro selection of a ribozyme corresponce of an oligonucleotide effector (AAAS7854). The Li ribozyme ligase was selected from a pool of RNA molecules comprising a central catalogues and a central catalogues and contral catalogues in length flanked on both sides by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                        Detecting an analyte in a sample comprises providing nucleic acid sequence which is catalytically active in presence of analyte, contacting catalytic nucleic acid with substrate and amplifying catalytic product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozyme; catalytic RNA; analyte detection; effector molecule; nucleic acid substrate; in vitro selection; ribozyme ligase; conformation dependent activity; allosteric activation; ss.
                                                                                                                                                                                                                                                           The invention relates to a method of detecting an analyte in a sample. The method comprises providing a nucleic acid sequence which is initially
                                                                                                                                                                                                                                                                                                               Disclosure; Page; 36pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                           Nathan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-OCT-1998;
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nilarity 73.1%;
Conservative
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/*tag=
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cc constant sequence regions (the N90 RNA pool; AAA57851). In the presence of the effector, selection was performed using one of the tagged substrate molecules AAA57855-A57857. RNAs with ligase activity (i.e., those which have become ligated to the substrate molecule) were reverse transcribed using the effector oligo, and then PCR amplified using the ceffector and a DNA primer identical in sequence to the substrate used for the selection. A ribozyme ligase, L1, was selected via this procedure. L1 can only adopt its active conformation (AAA57859) in the presence of the effector oligo (analyte). In the absence of the effector, L1 adopts an inactive conformation (AAA57860). The present sequence represents the deoxy-T22-tagged substrate oligonucleotide. The dT22 tag enables successfully ligated products to be isolated using oligo(dA) cellulose Type 7. Note: The present sequence is not given in the specification, but is created from the inforamtion given on page 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 19
                                                             Affixing a peptide acceptor to an RNA molecule useful for producing fusion proteins for isolating proteins or nucleic acids with desired properties through attachment of a peptide acceptor to the 3' end own molecule.
                                                                                                                                                         Kurz M,
                                                                                                                                                                                                               27-JUL-1999;
                                                                                                                                                                                                                                       19-JUL-2000; 2000WO-US019653
                                                                                                                                                                                                                                                                     01-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF60450;
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                                                                                                                                                                                     (PHYL-)
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/note= "A-TEG2"
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Pred. No. 2.
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The present invention relates to a method for affixing to an RNA molecule through the formation of a covalent

a peptide acceptor bond, noncovalent

Example

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Page

56pp;

English

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RESULT 3206

RAAL45359

AAL45359

AC AAL4535

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XX PAL4535

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  The present invention relates to the identification of specific proteolytically cleavable peptides by incubating a library of fusion molecules, comprising a peptide and nucleic acid encoding said peptide, with a proteolytically active sample, then isolating the cleavage tragments and determining the coding sequences in the separated fusion molecules. The coding sequences identified by the method are used for production of specifically proteolytically cleavable substances, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               stroke; neuronal disease; arthritis; pancreatitis; hypertension; thrombosis; infection; schistosomiasis; herbicide; insecticide; fungicide; cerebroprotective; neurological; antiarthritic; pancreatic; hypotensive; antithrombotic; virucide; protozoacide; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide cleavage; chemical active ingredient targeted release; diagnosis; antiasthmatic; osteopathic; cytostatic; asthma; osteoporosis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Puromycin linker
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                                                                                                                                                                                                                  Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                              Reimholz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-AUG-2000; 2000DE-01041238.
                                                                                                                                                                                                                                                                   Identifying specifically cleavable peptide, useful for targeted delivery and developing protease inhibitors, by incubating test with peptide-nucleic acid fusion.
                                                                                                                                                                                                                                                                                                                                                                                2002-269356/31.
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                                                                                                                                                                                                                    Page 19; 38pp; German.
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                          cc activity of cells comprising detecting, in a sample, the concentration cand/or subcellular localization of APC (denomatous polyposis coli) cc protein (I). The invention also describes (1) determining function of (I) cc in a sample by detecting presence of the C-terminal, DNA-binding domain of (I); (2) detecting presence of the C-terminal, DNA-binding domain of (I); (3) purifying, enriching and/or detecting ct the DNA-binding domain of (I); (3) purifying, enriching and/or detecting cc (I) or its fragments by reaction with a probe; (4) double-stranded DNA (CC (S2); (5) peptide fragment of (I) containing at least the DNA-binding cc domain; (6) antibodies (Ab) directed against an epitope of positions 1340 cc (1901, 2219-2580 or 2581-2843 of (I); (7) set of two or more antibodies, (CC (1-1299) of (I); and (8) kit for detecting DNA-binding capacity of (I) or its fragments in a sample consisting of (II), Ab or the set of (7). The method is used to detect proliferative, especially tumor (precursor), cells, to detect proliferative, especially tumor (precursor), and purify and to purify capacity of (I), and to purify capacity of (I) and to purify capacity of (II) and to purify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Determining proliferative capacity of cells, useful e.g. for detecting tumor cells, by measuring concentration and subcellular localization of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APC binding protein; cell proliferation; adenomatous polyposis coli; tumor cell detection; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 10; Page 13; 26pp; German.
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Pred. No. 2.7e+03
2; Mismatches
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ADC16682/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 19; Conserv
                                                           Query Match
Best Local
                                                Matches
                                                                                                                            The invention relates to a novel polynucleotide comprising a sequence encoding a ribozyme that can aminoacylate tRNA with a non-cognate amino acid. Ribozymes encoded by the polynucleotide of the invention are used to prepare polypeptides that contain non-cognate, including non-natural, amino acids. The invention more specifically provides catalytic RNA molecules having cis aminoacylation activity with a catalytic and aminoacylation domain, or an RNA molecule with trans aminoacylation activity with only a catalytic domain. The products of the invention are potentially useful for biomedical and therapeutic use, e.g. for probing the structure and function of proteins; preparation of peptide libraries and in proteomics. This polynucleotide sequence represents a DNA oligo relating to the RNA molecule with aminoacylation activity of the
                                                                                                                                                                                                                                                                                                                            New polynucleotide, useful for preparing peptides containing non-cognate amino acids, encodes ribozyme that can aminoacylate tRNA with such amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003070740-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  aminoacylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC16682
                                                                                                                                                                                                                                                                                            Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                            15-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-FEB-2003; 2003WO-US005007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aminoacylation RNA molecule related DNA oligo, P3-2.
                                                                                                                                                                                                                                                                                                                                                                                                                    ( YNYU)
                      4004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4018 AGAAAAAAGAGAGAAAACAAAATGTT 4043
26
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                                               19; Conser
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                                                                                                                                                                                                                                                                                                                                                                                          Murakami H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30
                     TTAGGTCTAAAATGAGAAAAAAGAGA 4029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aminoacylate; tRNA; non-cognate; catalytic
ation; trans; proteomic; ds.
                                                                                                                                                                                                                                                                                          SEQ ID NO 42; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%; ilarity 73.1%; Conservative
                                                                                            BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP;
                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            2002US-0357424P.
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                                                       0.2%;
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                                                                                            C; 1
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                                            Score 14.8; D
Pred. No. 2.9e
0; Mismatches
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Pred. No. 2.9e
0; Mismatches
                                                                                            21 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                    FOUND.
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                                                          .9e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .9e+03;
                                                                       DB 1;
                                                                                            0 Other;
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                                                                   Length 30;
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                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA molecule;
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                                             Gaps
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RESULT 3209

RESULT 3210
ABK32799/c
ID ABK32799/X
X ABK3279
XX ABK327

Human APPBP1 gene,

23-APR-2002 ABK32799

(first entry)

ABK32799 standard;

DNA;

15

ВР

0

WO200202820-A1

Human; amyloid beta precursor protein binding protein 1; APPBP1; probe; Alzheimer's disease; transgenic animal; platelet aggregation; single nucleotide polymorphism; SNP; allele-specific oligonucleotide; ss

allele-specific oligonucleotide #29

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ALL44170/c
ID ALL441
XX PALL441
XX PALL441
XX POIPHY
XX 
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                                                                                                                     Best Loc
Matches
                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                              The invention comprises a method for the production of cytochrome C. The method involves culturing a prokaryote that has been transformed with a vector encoding a signal peptide and a cytochrome C maturation protein. The method of the invention is useful for producing cytochrome C. Cytochrome C produced by the method of the invention is used in reagent; and drugs for trapping nitrogen oxide (e.g. in purifying polluted atmosphere by trapping nitrogen oxide). The present DNA sequence represents a Porphyra yezoensis cytochrome C - related PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Production of cytochrome c by culturing prokaryote transformed with vector containing e.g. DNA of signal peptide and of eukaryotic cytochrome c maturation protein for use in reagents and drugs for trapping nitrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL44170 standard; DNA;
                                                                                                                                                                                                                                                 Sequence 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polluted atmosphere
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytochrome C; ss; maturation protein; nitrogen oxide trapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Porphyra yezoensis cytochrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JAN-2001; 2001JP-00014510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL44170;
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                                                                                                                                                  Local
                                                      4018 AGAAAAAAGAGAGAAAACAAAATGTT 4043
30
                                                                                                                     . Similarity 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UNIV NIPPON
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AAAAAAAAAAAAAAAAAAAAGGAT
                                                                                                                                                                                                                                           BP; 1 A;
                                                                                                                     Conservative
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                                                                                                                                                                                                                                           3 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     purification; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                               0.2%;
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                                                                                                               Pred. No. 3e+00; Mismatches
                                                                                                                                                                                  Score 14.8;
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                                                                                                                                                                                                                                           26 T; 0 U; 0 Other;
                                                                                                                                               3e+03;
                                                                                                                                                                           DB 1;
                                                                                                                     7;
                                                                                                                                                                               Length 33;
                                                                                                                        Indels
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AAT66167/c
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for candidate drugs to treat diseases related to APPBP1 activity, in studying the effect of the variation on the biological activity of APPBP1, and the binding affinity of candidate drugs targeting APPBP1 for the treatment of disorders such as Alzheimer's disease. The haplotyping methods are useful in validating APPBP1 as a candidate target for treating a specific condition or disease predicted to be associated with APPBP1 activity, or in the design of clinical trials of candidate drugs for treating a specific condition or disease associated with APPBP1 activity. The transgenic animals are useful for studying expression of the APPBP1 isogenes in vivo, for in vivo screening and testing of drugs targeted against APPBP1 protein, and for testing the efficacy of therapeutic agents and compounds for disorders related to platelet aggregation in a biological system. ABK32771-ABK32337 represent human APPBP1 gene allele-specific oligonucleotides used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amyloid beta precursor protein binding protein 159 kD (APPBP1) gene polymorphic variants, useful e.g. in studying the expression and function of APPBP1 and screening candidate drugs for treating Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            which is a polymorphic variant of a reference sequence for the amyloid beta precursor protein binding protein 1, 59kD (APBBP1) protein or its fragment. The polymorphic variants are useful in studying the expression and function of APBBP1, in expressing APBBP1 protein for use in screening for candidate drugs to treat diseases related to APBBP1 activity, in
                                                                                                                                                            Glucosinolate; UDP-glucose:thiohydroximate S-glucosyltransferase; S-GT; transgenic plant; rapeseed oil; oilseed rape; canola; Brassica napus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 17; Page 13; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anastasio AE,
Stephens CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUN-2000; 2000US-0215511P
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               31-OCT-1995;
                                               07-MAY-1997.
                                                                                                                                           polymerase
                                                                                                                                                                                                          UDP-glucose:thiohydroximate S-glucosyltransferase primer
                                                                                                                                                                                                                                            15-JUL-1997
                                                                                                                                                                                                                                                                                                          AAT66167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                           TYTTTTTTTTTTG
                                                                                                                                           plant; rapeseed chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                          (first entry)
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               95EP-00402425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 1 C; 0 G;
                                                                                                                                                                                                                                                                                                          DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
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                                                                                                                                           PCR; primer; ss
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Pred. No. 1.5e
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 15
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                                                                                                                                                                                                            g13.
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RESULT 3212
ABN83985
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Degenerate primers based on 7 peptide sequences (AAW09826-32) of Brassica oleracea UDP-glucose:thiohydroximate S-glucosyltransferase (S-GT) were used in the PCR-RACE amplification of Brassica napus S-GT cDNA (see also AAT66166). Primer gI3 (AAT66167) was combined with the Anchor Primer of the Clontech 3'RACE kit (AAT66170), and the product was used as template in a second semi-nested PCR to yield S-GT partial clone pGL2-7 (AAT66173)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plants genetically transformed to interfere with glucose: thiohydroximate S-glucosyl:transferase ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 1 C; 1 G; 4 T; 0 U; 8 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for production or rapeseed oil with reduced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-247418/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Van Audenhove K, Pe
Hemmingsen SM, Reed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PLBZ )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-OCT-1995;
                                                                                                                                                                                                                                                                                                                             Leishmania parasite;
                                                                                                                                                                                                                                                                                                                                                     Leishmania
                                                                                                                                                                                                                                                                                                                                                                           02-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                         ABN83985 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2;
                                                                                         Molecular differentiation of Leishmaniosis parasites consists of detection by PCR RFLP technique via initiators and nucleotides.
                                                                                                                            WPI; 2002-417703/45.
                                                                                                                                                    Romanha
                                                                                                                                                                                                          28-SEP-2000; 2000BR-00004507.
                                                                                                                                                                                                                                 28-SEP-2000; 2000BR-00004507
                                                                                                                                                                                                                                                        30-APR-2002.
                                                                                                                                                                                                                                                                                BR200004507-A
                                                                                                                                                                                                                                                                                                       Leishmania
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                                                                                                                                                                         (-IMYU)
                                                                                                                                                                                     (FIOC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PLANT GENETIC SYSTEMS NAT RES COUNCIL CANADA.
                                                                                                                                                                         FIOCRUZ FUNDACAO CRUZ OSWALDO.
UNIV FEDERAL MINAS GERAIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
Similarity 55.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AARTTRAARTTNSWYTCYTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page
                                                                                                                                                                                                                                                                                                                                                    kinetoplast mini circle DNA PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                  Volpini
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peferoen M, Grootwassink JWD,
sed DW, Kolenovsky AD;
                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DW,
                                                                                                                                                  AC,
                                                                                                                                                                                                                                                                                                                              pharmaceutical; kinetoplast; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                   De Azeredo Passos VM,
                                                                                                                                                                                                                                                                                                                                                                                                                         ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.6; DB 1
Pred. No. 2.1e+03
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                                                                                                                                                                                                                                                                                                                               primer;
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The invention relates to the molecular differentiation of Leishmaniasis parasites. The method of the invention comprises detection of parasites of the Leishmania (Viannia) braziliensis, Leishmania (Viannia) guyanensi and Leishmania mazonensis species. The method employs initiators

buffer solution in a polymerase

2; Page 11; 38pp; Portuguese

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RESULT 3213
AAQ90391
ID AAQ9039
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Query Match
Best Local S
Matches 16
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Matches 14
                                                                             The sequences represented by, AAQ90390-90401 are synthetic DNA probes containing 3' ribonucleoside termini. The sequences shown in AAQ90402-15 are synthetic DNA probes with 5' amino termini. These sequences were specific for the polymorphisms of HIA gene dQa. The sequences were used in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CP-1;
SAED;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      restriction length polymorphism (PCR-RFLP) technique. Methods of the invention are useful in pharmaceuticals. The current sequence represents a PCR primer for amplification of conserved regions of kinetoplast mini
                                                                                                                                                                                                                                    New self-addressable electronic devices - used for multi-step multiplex reactions such as DNA hybridisation(s), clinical dia
                                                                                                                                                                                                                                                                                                                                             01-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              misc_feature
                                               Sequence
                                                                                                                                                                                                   Example 1; Page 40; 86pp; English.
                                                                                                                                                                                                                                                                                               Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                    26-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                             11-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                  WO9512808-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CP-1 (synthetic DNA probe with 3'ribonucleoside terminal #2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ90391;
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                                                                                                                                                                                                                                                                                                                        (NANO-) NANOGEN
             Local Similarity
                                                                                                                                                                                                                          bio:polymer synthesis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLA; dQa; 3' r: hybridisation;
 16;
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14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
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                                                                                                                                                                                                                                                                                               7
                                               BP;
 Conservative
                                                                      of these
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3' ribonucleoside; self-addressable electronic
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/note=
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           0.2%;
                                                                   reactions are greatly improved
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88
                                                                                                                                                                                                                                                                                                                                                                                                                                     "3' ribonucleoside terminal"
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Pred.
1; Mi
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                   Score 14.6;
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Pred. No. 2.1e+03;
                                             G; 0 T; 1 U; 0 Other;
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           2.3e+03;
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                        DB 1;
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 4.
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AAT10743
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Matches
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                                                                                                                    are used in the construction of the electronically self-addressable device (ED) of the invention. The ED comprises a substrate, an electro or opt. a number of electrodes supported by the substrate, a current source operatively connected to the electrode and an attachment layer adjacent to the electrode which is permeable to a counterion but not permeable to a molecule capable of insulating or binding to the electrode. The attachment layer is capable of attaching a macromolecul The ED is used for genetic typing and comprises a number of electronically addressable locations each comprising an electrode, and binding entity, such as one of these probes, attached to each of the locations capable of detecting the presence of a genetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4020 AAAAAAGAGAGAAAACAAAAT 4040
                                                                                               Sequence
                                                                                                                                                                                                                                                                                                             Electronically self-addressable e.g. nucleic acid hybridisation.
                                                                                                                                                                                                                                                                                                                                                                           Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Electronically self-addressable device; ED; electrode; current source; attachment layer; permeable; counterion; genetic typing; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT10743 standard; RNA;
                                                                                                                                                                                                                                                             The sequences given in AAT10742-67 are synthetic oligonucleotides which
                                                                                                                                                                                                                                                                                    Example 1; Page 60; 155pp; English
                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-097582/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               detection;
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                                                            Local
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                                                           Similarity
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AAAAAAAAAAAAAAAAAAA
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                                                                                                BP; 20
                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probe,
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/note= "3'-ribonucleoside terminus"
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                                                                                                                                                                                                                                                                                                                                                                           Evans
                                                          0.2%;
                                                                                               0 C;
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                                             Pred. No. 2.30
1; Mismatches
                                                           Score 14.6;
Pred. No. 2.
                                                                                               G; 0 T; 1 U;
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                                                                                                                                                                                                                                                                                                                          device -
                                                           2.3e+03;
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                                                                                                                                                                                                                                                                                                                          used
                                                                         DB 1;
                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                                          for electronic control of,
                                                                       Length 21;
                                                  Indels
                                                                                                                                                                       macromolecule.
                                                                                                                                                                                                                                    an electrode
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RESULT 3215
AAX81302
ID AAX8130
XX

AAX81302 standard; DNA;

21

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RESULT 3216
AAQ75780/c
ID AAQ7578
XX
AC AAQ7578
XX
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                              control multi-step and multiplex molecular biological reactions in microscopic formats. A key aspect of this inventions is played by the ion permeable permeation layer which overlies the electrode. This permeation layer allows attachment of nucleic acids to permit immobilization but also separates the attached oligonucleotides and hybridized target DNA sequences from the highly reactive electrochemical environment generated immediately at the electrode surface. The microelectronic device is designed and fabricated to actively carry out and control reactions such as nucleic acid hybridizations, antibody/antigen reactions, sample preparation, diagnostics and biopolymer synthesis. The device can electronically control the transport and attachment of specific binding entities, such as nucleic acids and polypeptides, to specific binding entities, such as nucleic acids and polypeptides, to specific micro-locations. The device can subsequently control the transport and reaction of analytes or reactants at the addressed specific micro-locations. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Microelectronic device; multi-step reaction; microscopic format; ion-permeable permeation layer; electrode; electrical control; transport; attachment; binding; DNA/RNA hybrid; probe; ss.
                   AAQ75780
                                                      AAQ75780 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                 specifically bound molecules, provide stringency control for DNA hybridization reactions and improve the detection of analytes. The present sequence represents a probe used to exemplify the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New microelectronic device designed to carry out and control multi-step and multiplex molecular biological reactions in microscopic format.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-385567/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sosnowski RG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3' ribonucleoside oligonucleotide probe CP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX81302;
                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes a self-addressable, self-assembling microelectronic device which is designed to actively carry out and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                       device is able to concentrate analytes and reactants, remove non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NANO-) NANOGEN INC.
                                                                                                                                                                              4020 AAAAAAGAGAGAAAACAAAAT 4040
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                                                                                                                                                                                                                l Similarity
16; Conser
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                                                                                                                                                                                                                                                                                  21 BP;
                                                                                                                                           AAAAAAAAAAAAAAAAAAA 21
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                                                                                                                                                                                                              0.2%;
larity 76.2%;
Conservative
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21
/*tag= a
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                                                                                                                                                                                                                                                                                  A; 0 C; 0 G; 0 T; 1 U;
                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
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                                                                                                                                                                                                             Score 14.6; DB 1; Length 21; Pred. No. 2.3e+03; Indels
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                                                                                                                                                                                                                                                                                   0 Other;
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RESULT 3217
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEO files AAO75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1995 (first entry)
                                                             16-APR-1993;
                                                                                                               01-NOV-1994.
                                                                                                                                       JP06303997-A.
                                                                                                                                                                                       aggregate;
                                                                                                                                                                                         Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                       04-AUG-1995
                                                                                                                                                                                                                                                                               AAQ75761;
                                                                                                                                                                                                                                                                                                         AAQ75761 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 9; 11pp; Japanese.
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          WPI; 1995-018287/03
                                   (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                     16-APR-1993;
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Pred. No. 2.3e+03;
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Matches 17
This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is action a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gen present in cells of the precancerous condition, where the normal somaticells of the patient are heterozygous for the first gene, the inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; call especific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                            Disclosure;
                                                                                                                                                                                          prevention dysplastic
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                                                                                                                                                                                     target genes for allele-specific drugs - used for diagnosis, and treatment of, e.g. cancers, atherosclerotic plaque, lesions, endometriosis or graft versus host disease.
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                                                                                                                    The oligonucleotide sequence is used as a probe to selectively detect Agrobacterium rhizogenes-derived plasmid DNA, as inserted into a plant genome. It is chemically synthesised and is complementary to T-DNA codifor the Ri plasmid of A.rhizogenes. It can also be used as a PCR primes and can easily detect a particular extraneous gene from a transformant plant with high sensitivity and selectivity
                                                                                                                                                                                                                                   Oligo:nucleotide for detection of transformant of plant chain reaction process to enable detection of specific with high sensitivity and selectivity.
                                                                                              Sequence
                                                                                                                                                                                                            Claim 1; Page 1;
                                                                                                                                                                                                                                                                                       WPI;
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Pred. No. 2.3e+03;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligo-nucleotide analogues which modulate arachidonic acid metabolism for treatment and diagnosis of conditions caused by lipoxygenase, phospholipase, leukotriene(s) etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             phosphorothioate linkage; SF-PLA2; ss.
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     JP05111490-A.
                                                                                                              Human leukocyte
                                                                                                                                                                HLA type analysis method
                                                                                                                                                                                                                      07-OCT-1993
                                                                                                                                                                                                                                                                                                                               AAQ42900 standard;
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                                                                                                           antigen; HLA
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Pred. No. 2.
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                                                                                                                method;
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Sequence 21

BP; 3 A;

7 C;

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0 Other;

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RESULT 3222
AAQ42902/c
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequence is that of DR1 & DR4 primer RRGb which was used a method of HLA type analysis involving amplification of a HLA cogene, or fragments of it, using 2 or more kinds of primers by polymerase method and subsequent restriction enzyme digestion analysis. The method enables easier analysis of HLA type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HLA type analysis method and its reagents - includes e.g. amplification of HLA class II gene, digestion by restriction enzyme, electrophoresis
                   The sequence is that of DR3, 5, 6, 8 primer RRGb which was used of a method of HLA type analysis involving amplification of a HI II gene, or fragments of it, using 2 or more kinds of primers by polymerase method and subsequent restriction enzyme digestion aranalysis. The method enables easier analysis of HLA type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 3 A; 7 C; 6 G; 5 T; 0 U;
                                                                                                                          HLA type analysis method and its reagents - includes e of HLA class II gene, digestion by restriction enzyme,
                                                                                                                                                                                                                                                        07-MAY-1993.
                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                           Human leukocyte
                                                                                                                                                                                                                                                                                                                                                 HLA type analysis method DR3,
                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ42902 standard;
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                                                                                         Example; Page 16; 21pp; Japanese.
                                                                                                                 and
                                                                                                                                                             WPI; 1993-184838/23.
                                                                                                                                                                                                          29-AUG-1991;
                                                                                                                                                                                                                               02-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SUMO)
                                                                                                                                                                                    (SUMQ ) SUMITOMO METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection.
                                                                                                                 detection.
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17; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUMITOMO METAL
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                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                           antigen; HLA
                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                    IND LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6; DB 1
Pred. No. 2.3e+03
                                                                                                                                                                                                                                                                                                                           classII
                                                                                                                                                                                                                                                                                                                                                 5
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                                                                                                                                                                                                                                                                                                                           gene; DNA polymerase
                                                                                                                                                                                                                                                                                                                                                  primer RRGb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                            includes e.g. amplification lon enzyme, electrophoresis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
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AAQ35258/c
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                                                                                                                                                         Query Match
Best Local :
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Best Local Similarity
Matches 17; Conserval
                                                                                                                                                   Matches
                                                                                                                                                                                          The sequence is that of a PCR primer which is used as part of a method for the detection of a gene relating to dwarfism of Agrobacterium rhizogenes. The method provides highly sensitive, easy and highly selective detection
                                                                                                                                                                                                                                        Oligo:nucleotide for detecting plant transforming principle coded to DNA of agrobacterium rhizogenes.
                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                      Agrobacterium
                                                                                                                                                                                                                                                                                                                                                                                    24-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                              AAQ35258 standard; DNA;
                                                                                                                                                                                                                           Claim 2;
                                                                                                                                                                                                                                                            WPI; 1993-030366/04.
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                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                          Polymerase
                                                                                                                                                                                                                                                                                                                                                                                                 AAQ35258;
                                                                                                                                                                                                                                                                          (SHMA ) SHIMADZU CORP
                                                                                                                                3007 CTCACCCCATCTTGTCACATC 3027
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1610
                                                                                                                         21
                                                                                                                                                0.2%;
11 Similarity 81.0%;
17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                             21 BP; 7
                                                                                                                                                                                                                           Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGAACTTCACAGACCAGCTGC 1630
                                                                                                                                                                                                                                                                                                                                                                                                                                               AGAGCTTCACAGTGCAGCGGC
                                                                                                                         CTCATCGCTGCTTGTCACATC 1
                                                                                                                                                                                                                                                                                                                                                          chain reaction; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                          2; 10pp; Japanese.
                                               rhizogenes dwarfism
                                                                                                                                                                                                                                                                                                                                                                      rhizogenes dwarfism
                                                                                                                                                                                                                                                                                       91JP-00128924
                                                                                                                                                                                                                                                                                                     91JP-00128924.
                                  reaction; detection;
                                                                                                                                                                            A; 3 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
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                                                                                                                                                        Score 14.6; DB 1;
Pred. No. 2.3e+03;
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Pred. No. 2.3e+03;
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                                                PCR
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RESULT 3225
AAQ35268/c
ID AAQ35268 standard; DNA;
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Best Local
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Best Local
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                                                                                                                                                  Oligo:nucleotide for detecting plant transforming coded to DNA of agrobacterium rhizogenes.
                                      Sequence 21
                                                                       The sequence is that of a PCR primer which is used as part of a method for the detection of a gene relating to dwarfism of Agrobacterium rhizogenes. The method provides highly sensitive, easy and highly
                                                                                                                           Claim 2; Page 2; 10pp; Japanese.
                                                                                                                                                                                        WPI; 1993-030366/04.
                                                                                                                                                                                                                                         31-MAY-1991;
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                                                                                                                                                                                                                                                                                                                     JP04356189-A
                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                      Polymerase
                                                                                                                                                                                                                                                                                                                                                                                            Agrobacterium rhizogenes dwarfism gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ35268;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence is that of a PCR primer which is used as part of a motor the detection of a gene relating to dwarfism of Agrobacterium rhizogenes. The method provides highly sensitive, easy and highly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 2; 10pp; Japanese.
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                                                                                                                                                                                                                 (SHMA ) SHIMADZU CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 CTCATCGCTGCTTGTCACATC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 detection
                                                                                                                                                                                                                                                                                                                                                                      chain
                                    BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                          91JP-00128924
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                                                                                                                                                                                                                                                                                                                                                                      reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                    3 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          쁌
                                                                                                                                                                                                                                                                                                                                                                      detection;
                                    G; 3
Score 14.6; DB 1;
Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.6;
Pred. No. 2.
                                    T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 U;
                                    u;
                                                                                                                                                                                                                                                                                                                                                                                               PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                                                                                                                               primer.
           Length
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                                                                                                                                                               principle
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              21;
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Local

l Similarity 17; Conserv

Conservative

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Mismatches

Indels

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Gaps

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RESULT 3226
AAQ70288/c
ID AAQ7028
XX AAQ7028
XX AAQ7028
XX 25-MAR-
DT 13-APR-
XX Capture
KW Capture
KW Capture
KW nucleic
XX Synthet
XX PR27015
XX 02-SEP-
XX 02-FEB-
XX (INMR)
XX (INMR)
XX WPI; 1:
XX WPI; 1:
XX Solid-|
PT comprii
CC 165 rB|
CC 165 rB|
CC 165 rB|
CC 165 rC|
C
RESULT 3227
AAQ61922
ID AAQ6192
XX
AC AAQ6192
XC AAQ619
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The detection probe CT3 (AAQ70288) is complementary to a region of the 16S rRNA sequence of Chlamydia trachomatis which forms part of a hairpin loop. The probe is only able to hybridise to the target RNA in the presence of a capture probe (i.e. dCT4, AAQ70289) immobilised on a support. The capture probe destabilises the secondary structure of the hairpin loop. The oligonucleotide probes CT3 and dCT4 are useful for distinguishing Chlamydia trachomatis from other closely related Chlamydia species. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Solid-phase capture probes for nucleic acid hybridisation assays -comprising immobilised oligo:nucleotide capable of destabilisation secondary structure formed in target nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Capture probe;
nucleic acid hy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 18; Page 15; 21pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mabilat C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-SEP-1994
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     Inhibition; replication; herpes simplex virus; HSV;
                                                          Human type II phospholipase A2 inhibiting oligomer,
                                                                                                                   25-MAR-2003
04-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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13-APR-1995
                                                                                                                                                                                                     AAQ61922;
                                                                                                                                                                                                                                                             AAQ61922 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                6938
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                                                                                                                                                                                                                                                                                                                                                                                                           TATTTGGGCATCCGAGTAACG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGTTTGGGCATCCAAGAAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e; detection probe; 16S ribosomal RNA; Chlamydia;
hybridisation assay; ss.
                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93FR-00002127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.6; DB 1;
Pred. No. 2.3e+03;
0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                6958
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
  HIV; aging;
                                                             ISIS no 3192.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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     modified_base
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human cytomegalovirus; influenza virus; inflammation; telomere length; neurological disorders; phospholipase A2 activity; hyperproliferation; malignancy; cardiovascular disease; snake bite; malignancy; retard; ss
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Synthetic.

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misc_feature
Location/Qualifiers
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"Phosphorothioate intersugar linkages"

WO9408053-A1

14-APR-1994.

29-SEP-1993; 93WO-US009297

29-SEP-1992; 92US-00954185

(ISIS-) ISIS PHARM INC

Hanecak RC, Anders Ecker DJ, Vickers Anderson KP, Bennet ckers TA, Wyatt JR, Bennett ÇĘ, Imbach Chiang Į, Σ, Brown-Driver

WPI; 1994-135613/16.

New modified oligo-nucleotide contg guanine quartet - inhibits activity of viruses, e.g. HIV, and phospholipase A2 and modulates telomere length of chromosomes.

Disclosure; Page 26; 144pp; English.

The sequences given in AAQ61917-55 are oligonucleotides which contain a G4 stretch and which may be used for inhibiting phospholipase A2 enzyme activity. Oligonucleotides such as these may also be used for inhibiting activity of HSV, HIV, human cytomegalovirus or influenza virus, or for treating inflammatory and neurological disorders caused by phospholipase A2 activity in cases of hyperproliferation, malignancy, cardiovascular disease and snake bite. They may also be used for inhibiting division of malignant cells by modualting telomere length, which may also retard aging. (Updated on 25-MAR-2003 to correct PN field.)

Sequence 21 BP; σ A 2 C; 10 G; 3 T; 0 U; 0 Other;

Query Match Best Local S Matches 17 l Similarity 17; Conserv Conservative 0.2%; 0 Score 14.6; Pred. No. 2 Mismatches .3e+03; DB 1; Length 4. Indels 21; 0 Gaps 0

1631 GGAAGATTTCCAAGGATGCGG 1651 GGAAGGTTTCCAGGGAAGAGG 21

AAQ61989 standard; DNA; 21 ВP

AAQ61989;

25-MAR-2003 04-NOV-1994 (revised) (first entry)

HIV replication inhibiting oligomer, T13G4T4.

Synthetic Inhibition; replication; herpes simplex virus; HSV; HIV; retard; human cytomegalovirus; influenza virus; influenzation; telomera human cytomegalovirus; influenza virus; influenzation; hyperprolifez neurological disorders; phospholipase A2 activity; hyperprolifez malignancy; cardiovascular disease; snake bite; malignancy; agin hyperproliferation; lignancy; aging; ss. length;

Location/Qualifiers

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                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 3229.
AAT11995/c
                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence may be used for inhibiting replication. This sequence may be used deficiency virus (HIV). Oligonucleotides such as this may also be used for inhibiting activity of HSV, human cytomegalovirus or influenza virus, or for treating inflammatory and neurological disorders caused by phospholipase A2 activity in cases of hyper- proliferation, malignancy, cardiovascular disease and snake bite. They may also be used for inhibiting division of malignant cells by modualting telomere length, which may also retard aging. (Updated on 25-MAR-2003 to correct PN
19-NOV-1992;
                                                                                                                                                                                                     CMV antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hanecak RC, Anuc...
NT. Vickers
                                                                                                                                                                   antisense; cytomegalovirus;
intermediate early complex;
                                                                                                                                                                                                                            25-MAR-2003
13-MAR-1996
                                                                                                                                                                                                                                                                                     AAT11995 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New modified oligo-nucleotide contg guanine quartet - inhibits act of viruses, e.g. HIV, and phospholipase A2 and modulates telomere
                                                                                                            modified_base
                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                               AAT11995;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 21; Page 60; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of chromosomes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-APR-1994.
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                     25-JAN-1993;
                                                                   US5442049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1994-135613/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-)
                                                                                                                                                                                                                                                                                                                                                         4468 TITITITITITITITITGTCTT 4488
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SISI
                                                                                                                                                                                                                                                                                                                                            TTTTTTTTTTGGGGTTTT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 0 A; 0 C; 4 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anderson KP, Bennett CF, Chian ickers TA, Wyatt JR, Imbach JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                                      oligonucleotide
9208-00927506
                     9308-00009263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92US-00954185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93WO-US009297
                                                                                                            Location/Qualifiers
                                                                                       'note=
                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Labeled with 32P"
                                                                                       "phosphorothioate backbone"
                                                                                                                                                                    CMV;
                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.6;
Pred. No. 2
                                                                                                                                                                                                      (ISIS 4847).
                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                   human; therapy; prophylaxis; diagnosis; IE2; DNA polymerase gene; ss.
                                                                                                                                                                 DNA polymerase gene;
                                                                                                                                                                                                                                                                                                                                                                                                  2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chiang M,
                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brown-Driver VL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inhibits activity
                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          length
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RESULT 3230
AAT01661/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A series of 21 phosphorothioate antisense oligonucleotides (ONs) (AATI1987- 2007) were examined for anti-cytomegalovirus (CMV) activity. ISIS 4847 targets the DNA polymerase gene 5' untranslated region. It has an IC50 value of 0.9 microM. (ONs which inhibit CMV at one-third the dosage (or below) at which the negative control shows activity in this experimernt (IC50 = 1 microM. or less) are preferred). Antisense ONs targeting CMV DNA or RNA coding for the IE1, IE2 or DNA polymerase proteins have been shown to be effective in therapy, prophylaxis and diagnosis of CMV infection. The ONs may be modified to reduce nuclease resistance and to increase their efficacy. Modifications include phosphorothioate backbones, alkyl and halogen-substituted sugar moieties at the 2' position. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligo-nucleotide inhibits cytomegalovirus replication - by binding to a portion of cytomegalovirus RNA, for the diagnosis, prophylaxis and treatment of CMV diseases.
WPI; 1995-090841/12.
                       Anderson KP,
                                                                            09-AUG-1993;
                                                                                                     09-AUG-1994;
                                                                                                                                                         WO9504748-A1
                                                                                                                                                                                                                                                   misc_feature
                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                     antiviral;
                                                                                                                                                                                                                                                                                                                                 peptide nucleic acid;
                                                                                                                                                                                                                                                                                                                                                          Peptide nucleic acid targetting CMV DNA pol 5'-UTR
                                                                                                                                                                                                                                                                                                                                                                                                                AAT01661;
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT01661 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Baker B,
                                                                                                                                                                                                                                                                                                                                                                                      17-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3024 CATCTGGCCCTGACCCCACTG 3044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
17; Conser
                                                 SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12; Col 17-18; 66pp; English.
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                                                                                                                                                                                                                                                                                                                     diagnostic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                   PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                       Crooke ST,
                                                                            93US-00104438
                                                                                                      94WO-US009039
                                                                                                                                                                                 /note "at least one (and preferably all) of the backbone subunits are composed of amide units, so that the oligomer consists of the nucleobases attached covalently to a polyamide backbone"
                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 C; 9 G; 0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anderson
                                                                                                                                                                                                                                                                                                                     PNA;
                        Mirabelli
                                                                                                                                                                                                                                                                                                                                 cytomegalovirus; CMV; papillomavirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.6; DB 1;
Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 U;
                       Ś
                       Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                        Cowsert LM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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New peptide nucleic acid oligomers hybridisable to cytomegalovirus or papilloma:virus - are stable anti:sense molecules with high affinity single stranded DNA, used for treating infections.

Claim 2;

Page 44; 65pp; English

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RESULT 3231
AAT35317/c
ID AAT35311/X
XX AAT3531
XX AAT3531
XX AAT3531
XX 25-MAR-
DT 25-MAR-
DT 13-NOV-
XX 13-NOV-
XX PTP; px
KW Chromos
KW Chro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New oligomers are claimed which (A) have at least one peptide nucleic CC acid (PNA) subunit and (B) have a sequence hybridisable to AUG region, 5' CC untranslated region, intron/exon junction or coding sequence of CC cytomegalovirus gene selected from DNA polymerase, IE1 and IE2, or CC hybridisable to the E, E2, E4, E5, E6, E7, L1 or L2 reading frames of a CC papillomavirus. The PNAs can be used to target RNA and single stranded CC DNA (ssDNA) to produce antisense-type gene regulation moieties. Hence they may be used therapeutically for modulating cytomegalovirus and CC papillomavirus processes and also as diagnostics (e.g., as probes for CC specific mRNAs). PNA oligomers have high affinity for complementary CC single stranded DNA. They are also able to form triple helices in which a first pNA strand binds with RNA or ssDNA and a second pNA strand binds CC with the resulting double helix or with the first pNA strand. The PNAs CC possess no significant charge and are water soluble, which facilitates CC cellular uptake. Futher, since they contain amides of non-biological amino acids, they are biostable and resistant to enzymatic degradation by croteases. The present sequence targets CMV DNA polymerase 5' CC untranslated region (5' UTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                            26-JUN-1991;
31-JAN-1992;
01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chromosome 12p; abnormality; wuvarion, ---- regulation; cancer; leukaemia; diagnosis; megakaryocyte regulation; nolymerase chain reaction; ds.
      Detecting 12p chromosomal abnormality associated with neoplastic disease
                                                                 WPI; 1996-341506/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SH-PTP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003
13-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT35317 standard; cDNA; 21 BP
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                                                                                                                                                                                        (BETH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5536636-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PTP; protein tyrosine phosphatase; SH2; Src homology region 2; chromosome 12p; abnormality; mutation; detection; probe; neoplasia;
                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3024 CATCTGGCCCTGACCCCACTG 3044
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l Similarity 81.0%;
17; Conservative
                                                                                                                                                                                        BETH ISRAEL HOMESSACHUSETTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          criciédecerédecedencie 1
                                                                                                                      Rosenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6 A;
                                                                                                                                                                                        ISRAEL HOSPITAL ASSOC.
ACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene PCR primer,
                                                                                                                                                                                                                                                                         91US-00721112.
92US-00829141.
92US-00983926.
                                                                                                                                                                                                                                                                                                                                                                                                  94US-00202389
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                                                                                                                            80,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 C; 9 G; 0 T; 0 U; 0 Other;
                                                                                                                            Freeman RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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Pred. No. 2.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            400B.
                                                                                                                         Plutzky J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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Example 3; Col 8; 14pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 3232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC human cDNA sequence encoding SH-PTP1 (protein tyrosine phosphatase-1, CC with two SH2 domains). The primers are used in the analysis of the SH-CC PTP1 gene of patients with acute lymphoblastic leukaemia (ALL). A CC fragment complemetary to the SHPTP-1 sequence from nucleotides 537-653 is CC domains of the two tandems of the SHPTP-1 sequence from nucleotides 537-653 is CC domains for the two tandems SH2 domains (see AAR99312). If the probe cc region coding for the two tandems SH2 domains (see AAR99312). If the probe CC differentive of an abnormality, normally associated with neoplastic disease, in partic, ALL. The probe hybridises to a part of the CC differentive of an abnormality, normally associated with neoplasia.

CC Alternatively the wild-type SH-PTP1 or SH-PTP2 gene or protein may be comparison to sequenced PTP genes taken from a patient, where CC differences indicate an abnormality. The activity of SH-PTP1 may also be composely altered by mutation to effect a change in megakaryocyte content of the first production and hence platelet production. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV; infection inhibitor; triplex forming; purine rich promoter; V3 loop; transcription inhibitor; gp120 protein; viral growth; enzyme inhibitor; PLA2; telomere length; glove coating; condom; 88.
                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV inhibitor #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
03-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 4 A; 10 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               function and correct PF fi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Fig 9; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT35001;
                                           New octa:nucleotide with guanosine quartet and phosphoro:thioate links is inhibitor of HIV infection by binding to the V3 loop.
                                                                                            WPI; 1996-285782/29
                                                                                                                            Imbach JL,
                                                                                                                                                                                            29-SEP-1992;
                                                                                                                                                                                                                           28-SEP-1993;
                                                                                                                                                                                                                                                           04-JUN-1996.
                                                                                                                                                                                                                                                                                           US5523389-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT35001 standard;
                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         using SH-PTP1 protein tyrosine phosphatase gene specific probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4436 CTAGGGCATGTGGGTGGGTGG 4456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 CAAGGTCATGTGCGAGGGTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                field.)
                                                                                                                            Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                                                                                                                                                                                                            93US-00128011
                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                         /note= "phosphorothioate nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                            Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.6; DB 1
Pred. No. 2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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RESULT 3233
AAT31784/c
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Best Local S
Matches 17
Determining whether a human patient has a tumour or a metastasised tumour, comprises determining whether a cytokeratin 20 (CK20) gene prod. is present in a tissue sample that does not normally contain CK20. This method is partic. useful for the detection of epithelial cell tumours, e.g. colorectal, stomach, mucinous ovarian or gall bladder adenocarcinoma or bladder or transitional cell carcinoma. In an example RNA was extracted from normal human blood samples and samples spiked with adenocarcinoma HT29 cells, transitional cell carcinoma RT112 cells or breast adenocarcinoma MCF-7 cells and subjected to PCR amplification using the CK20, CK8 or CK19 mRNA specific primers AAT31769/70, CNB and CK19 mRNA was not detected in normal blood, but a respective 370, 244 or 214 bp prod.
                                                                                                                                                                                                                                                                                                                                                                                  Detection of gene prod.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tresidues (such as these sequences) are triplex forming oligonucleotides, and form purine rich promoter elements used to inhibit transcription. These sequences bind to the HIV 99120 protein at the V3 loop via the internal guanosine quartet. This binding prevents cell-to-cell and virus-to-cell infection. The sequences may also be used for inhibiting viral growth, and other viral genes, for inhibiting the enzyme PLA2, and to modulate telomere length. In some cases these sequences need to be chemically modified. The chemically modified oligonucleotides preferably include at least one phosphorothioate linkage. Other modified intersugar links, or 2'-modified sugar residues can also be used. These oligonucleotides can be used for coating gloves, condoms, etc., or for topical application. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Selby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9617080-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            epithelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Determination;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytokeratin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    784/c
AAT31784 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT34998-T35001 represent HIV inhibitors. Sequences containing only G and T residues (such as these sequences) are triplex forming
                                                                                                                                                                                                                                                                                                                                     Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-277793/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                epithelial cell; colorectal; stomach; adenocarcinoma; bladder; transitional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT31784;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (IMCR ) IMPERIAL CANCER RES TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ę,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTTTTTTTTTTTTTTGTCTT 4488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 BP; 0 A; 0 C; 4 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTTTTTTTTTTTGGGGTTTT
                                                                                                                                                                                                                                                                                                                                                                                                               of
f
                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Burchill SA
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                                                                                                                                                                                                                                                                                                                                                                                    ın a
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                                                                                                                                                                                                                                                                                                                                                                                  human tumours or metastasis
n a tissue sample which does
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94GB-00023912.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95WO-GB002734.
                                                                                                                                                                                                                                                                                                                                   17; 36pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     umour; metastasis; cytokeratin 19;
colorectal; stomach; mucinous ovar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
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                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mucinous ovarian; gall
cell; carcinoma; primes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                       not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                  by detecting a cyto-keratin ot normally contain CK20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             carcinoma; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CK19; detection; rian; gall bladder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 3234
                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                               This sequence represents an initialising oligonucleotide for a fragment of gUC19. The invention relates to a method for the identification of a sequence of nucleotides in a polynucleotide (PNT), which involves: (a) extending an initialising oligonucleotide (ONT) along the PNT by ligating an ONT probe to form a duplex; (b) identifying one or more nucleotides of the PNT; and (c) repeating steps (a) and (b) until the nucleotide sequence is determined. The method is useful e.g. in gene function and control investigations, genomic analysis, genetic identification, forensic analysis, genetic counselling or medical diagnostics. The method avoids the need for electrophoratic separation of similarly sized DNA fragments, eliminates the difficulties associated with detection and analysis of spatially overlapping bands of DNA fragments in a gel or similar medium and avoids the need to generate DNA fragments from long
                                                                                                                                                                                                                                                                                                                                  Sequencing DNA by parallel oligo:nucleotide extension - extending initialising oligonucleotide by ligating probe avoiding electrophoretic sepns.
                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                     single-stranded
                                                                                                                                                                                                                                                                                                           Example 1; Page 20; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Initialising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX33889;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX33889 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    was detected in the
                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      initialising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 forensic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                       (SPEC-)
                       3441
                                                                                                                                                                                                                                                                                                                                                                                      1996-485723/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   985
21
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                                                                                                                                                                                                                                                                                                                                                                                                                                       SPECTRAGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence determination; genomic analysis; genetic identification;
c analysis; genetic counselling; medical diagnostic;
                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
               CCCCACCTTACTTCTCCTCCC 3461
                                                                                                                                                                                                                                                                                                                                                                                                              S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATGCAGATCGAAGGCCTGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGGAGATCAAGGGCCTGAAG 1005
                                                                                               B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3
                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96WO-US005245
                                                                                                                                                                                                                                                                                                                                                                                                                                                             95US-00424663.
                                                                                                7
                                                                                                                      template with a DNA
                                                                                              ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                              0 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HT29, RT112 and MCF-7 spiked blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 C; 4 G;
                                                           81.0%;
                                                                      0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                              14 G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                           Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for pUC19 fragment amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 T;
                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 U;
                                                                                                                      polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .3e+03;
                                                           .3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                        DB 1;
                                                                                                0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.
                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                               probe
                                                                                                                                                                                                                                                                                                                                                             involving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                                                                                                                                                ő
                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                               form duplex,
                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                               0
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RESULT 3236
AAT86582
ID AAT8658
XX
AC AAT8658
XX
DT 25-MAR-
XX
DE Phospho
                                                                                                                                                                                                              RESULT 3235
AAX33890/c
                                                                                                                         밁
                                                                                                                                           Ş
                                                                                                                                                                  Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                           This sequence represents an initialising oligonuclectide for a fragment of pUC19. The invention relates to a method for the identification of a sequence of nucleotides in a polynucleotide (PNT), which involves: (a) sequence of nucleotides in a polynucleotide (PNT), which involves: (a) extending an initialising oligonucleotide (ONT) along the PNT by ligating an ONT probe to form a duplex; (b) identifying one or more nucleotides of the PNT; and (c) repeating steps (a) and (b) until the nucleotide sequence is determined. The method is useful e.g. in gene function and control investigations, genomic analysis, genetic identification, forensic analysis, genetic counselling or medical diagnostics. The method avoids the need for electrophoretic separation of similarly sized DNA fragments, eliminates the difficulties associated with detection and analysis of spatially overlapping bands of DNA fragments in a gel or similar medium and avoids the need to generate DNA fragments from long
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequencing DNA by parallel oligo:nucleotide extension - involving extending initialising oligonucleotide by ligating probe to form duplex,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer; sequence determination; genomic analysis; genetic i
forensic analysis; genetic counselling; medical diagnostic;
initialising oligonucleotide; amplification; ss.
 Phosphorothicate oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 20; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-485723/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-OCT-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Initialising oligonucleotide for pUC19 fragment amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX33890 standard; DNA;
                       25-MAR-1998
                                                                 AAT86582 standard;
                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                   single-stranded template with a DNA polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9633205-A1
                                                                                                                                              3442
                                                                                                                        21
                                                                                                                                                                  l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SPECTRAGEN INC.
                                                                                                                                         CCCACCTTACTTCTCCTCCCT 3462
                                                                                                                                                                                                                                                                                                                                                                                                                           electrophoretic sepns.
                                                                                                                         CICICCITCCCICICCICCI 1
                                                                                                                                                                                                              21
                                                                                                                                                                                                             BP; 8
                                                                                                                                                                 0.2%;
llarity 81.0%;
Conservative
                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95US-00424663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96WO-US005245
                                                                                                                                                                                                              A; 0 C; 13 G; 0 T; 0 U;
                                                                  DNA;
                                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21
                                                                  BP
                                                                                                                                                                             Score 14.6; DB 1; Length 21; Pred. No. 2.3e+03;
                                                                                                                                                                     Mismatches
                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genetic identification;
                                                                                                                                                                     Indels
                                                                                                                                                                   0,
                                                                                                                                                                     Gaps
                                                                                                                                                                   0
    RESULT 3237
                                                                                                                                                                                                                                                밁
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Matches
             12-JUN-1997
                                   WO9720929-A1
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Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a phosphorothioate oligonucleotide which was prepared by solid phase synthesis. The method comprises adding at least one dimeric phosphoramidite synthon, optionally having a protected thiosester group in its internucleotide link, during the synthesis cycle. These novel dimeric phosphoramidite synthons are used as antisense molecules for inhibition of gene expression. The method gives increased yields of the phosphorothiate oligonucleotide (since fewer cycles are needed) and facilitates separation of impurities (greater difference in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Solid phase synthesis of phosphoro:thioate oligo:nucleotide(s) using new dimeric synthon(s) - useful as anti:sense molecules for inhibiting gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 0 A; 10 C; 0 G; 11 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      size compared with use of monomeric synthons)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phosphorothicate oligonucleotide; dimeric phosphoramidite thicester; DNA synthesis; antisense oligonucleotide; gene
Synthetic.
                                                 Fibroblast growth factor; wound healing; cartilage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; Page 20; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-415290/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9729116-A1.
                                                                                                                   Rat fibroblast growth factor FGF-10 RACE primer
                                                                                                                                                                    12-FEB-1998
                                                                                                                                                                                                                   AAT77232;
                                                                                                                                                                                                                                                                 AAT77232 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CRUA-) CRUACHEM LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-AUG-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                 5327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CB,
                                            healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                       TCTCTCTTTGCCTCACTCTCT 5347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rao MV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
ilarity 81.0%;
Conservative
                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96GB-00002326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-GB000327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "Phosphorothioate linkages between alternate nucleotides (1 and 2, 3 and 4 etc.)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
1. .21
                                                                                                                                                                                                                                                                 21
                                                 rat; human; recombinant DNA; bone RACE primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthon;
therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
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                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
                  09-MAY-1997;
14-AUG-1997;
14-AUG-1997;
14-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-DEC-1995;
28-MAR-1996;
24-JUL-1996;
                                                       18-APR-1997;
25-APR-1997;
06-MAY-1997;
                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a RACE primer involved in the amplification of rat fibroblast growth factor FGF-10. Recombinant FGF-10, vectors, containing the DNA, and host cells, containing the vectors, are useful for the recombinant production of FGF-10. The recombinant FGF-10 is useful for the treatment of diseases and injury of bone or cartilage,
                                                                                                       01-OCT-1997;
                                                                                                                                           GB2317891-A
                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                   Human; telomerase reverse transcriptase; hTRT; TRT; diagnosis; prognosis; cell proliferation; cancer; ageing; ribonucleoprotein; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Recombinant fibroblast growth the treatment of bone disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-DEC-1996;
                                                                                    01-OCT-1996;
                                                                                                                          08-APR-1998
                                                                                                                                                                                                                                           primer; 88.
                                                                                                                                                                                                                                                                                 Telomerase
                                                                                                                                                                                                                                                                                                    13-AUG-1998
                                                                                                                                                                                                                                                                                                                                           AAV30731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-319776/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Itoh N,
(GERO-) GERON
                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SUMU ) SUMITOMO PHARM CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                            5740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as a wound healing promoter
                                                                                                                                                                                                                                                                                                                                                                                          ب
                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     TCCCTTTTCTTCTATTCACTC
                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 BP; 2 A; 9 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Negoro
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                                                                                                                                                                                                                                                                                 reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 35; 51pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                    (first
CORP.
                 96US-00724643.
97US-00844419.
97US-00846017.
97US-00851843.
97US-00854050.
97US-00911312.
97US-00912951.
97US-00915503.
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96JP-00103240.
                                                                                                       97GB-00020890.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96JP-00214378.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96WO-JP003579
                                                                                                                                                                /*tag=
/note=
                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                transcriptase primer 260-280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Katsumata
                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            factor FGF-10 and for wound
                                                                                                                                                                                                                                                                                                                                                                                                           5760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tagashira
                                                                                                                                                                                                                                                                                                                                                                                                                                      .3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 몂
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and related DNA - useful healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 1;
                                                                                                                                                                                                                                                                                                                                                                                                                              4.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for
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                                                                                                                                                                                                                              RESULT 3239
AAV08273/c
                                                                                                                                                                                                                                                      describes human telomerase reverse transcriptase (hTRT). The present clinvention also describes the following methods: (A) determining whether a cream compound is a modulator of hTRT, by detecting the change in hTRT recombinant protein or polynucleotide, on administration of the compound; (C) preparation of hTRT with a telomerase by contacting a protein of the complex formed and detection, and detecting the product and correlating the presence of RNA detection, and the product and correlating the presence of complex or amplification, and (B) the use of an agent that causes an increasing hTRT expression, and (B) the use of an agent that causes an increase in cell correspicate cell proliferation to create a medicament that inhibits ageing. A protein preparation of hTRT and the polynucleotide encoding hTRT can be used in the manufacture of medicament that inhibits to treat conditions that are associated with high telomerase activity can be used to treat conditions that are associated with high telomerase activity. A protein preparation of hTRT can also be used in the new methods
                                                                                                                                                           Matches
                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a primer from the present invention which describes human telomerase reverse transcriptase (hTRT). The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pure and recombinant human Telomerase Reverse Transcriptase and variants - are useful in the diagnosis, prognosis and treatment proliferation conditions especially cancer and ageing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
AAV08273 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYTE-) UNIV TECHNOLOGY CORP
                                                                                                                  4735 GGCCAGCTGGAGGAAGAAGGG 4755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TR,
                                                                                                                                                       l Similarity
17; Conserv
                                                                                       -ر
                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lingner
                                                                                                                                                                                                                              BP; 5
                                                                                                                                                         Conservative
                                                                                                                                                                                                                              P
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                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                            4 C; 11 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nakamura
21
ВP
                                                                                                                                                   0;
                                                                                                                                                                         Score 14.6; DB 1
Pred. No. 2.3e+03
                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 H,
                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chapman KB,
                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Morin
                                                                                                                                                                                           Length
                                                                                                                                                           Indels
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                                                                                                                                                         0
                                                                                                                                                         Gaps
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(BAYU)
(UYJO)
(UTAH)
                                                                                                                                     PCR primer;
                                                                                                                                              ATP binding cassette; A Fundus Flavimaculatus;
                                                                                                                                                                                                           AAV08273;
                                          27-FEB-1997;
                                                          27-FEB-1998;
                                                                            03-SEP-1998.
                                                                                            WO9837764-A1
                                                                                                              Homo
                                                                                                                      Synthetic
                                                                                                                                                                        PCR primer ABCR.EXON43:F for ABCR coding sequence
                                                                                                                                                                                          27-JAN-1999
                                                                                                             sapiens
BAYLOR COLLEGE MEDICINE.
UNIV JOHNS HOPKINS.
US DEPT HEALTH & HUMAN S
UNIV UTAH.
                                                                                                                                                                                         (first entry)
                                                           98WO-US003895
                                          97US-0039388P
                                                                                                                                                       ABC transporter; ABCR; Stargardt Disease; therapy;
                                                                                                                                               age-related macular degeneration; diagnosis;
         & HUMAN SERVICES
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XXCCCCXXXXPPPXXXPIIIXX

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RESULT 32
AAVO8280
ID AAVO8280
AC AAVO
XX AAVO
XX AAVO
XX AFP
CR XX AFP
KW FURN
XX SYNI
OS HOMM
PN WO91
XX OS HOMM
PN CUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local
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Lupski JR,
Sun H;
                                                                                                                                                                                                                                    Allikmets F
Lupski JR,
                                                                  Retina-specific ATP-binding cassette transporter and DNA -e.g. diagnosis and treatment of macular degeneration, such Stargardt Disease, Fundus Flavimaculatus and age-related de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a PCR primer for DNA encoding the human retina specific ATP binding cassette transporter (ABCR) of the invention. ABCR may be used in compositions for screening agents that alters ABCR. The agent can inhibit Stargardt Disease, Fundus Flavimaculatus and ageralated macular degeneration (MD). Primers (such as this sequence) and probes for the ABCR DNA can be used in a diagnostic kit for detecting M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Retina-specific ATP-binding cassette transporter and DNA -e.g. diagnosis and treatment of macular degeneration, such Stargardt Disease, Fundus Flavimaculatus and age-related de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9837764-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV08280 standard; DNA; 21 BP
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                                         An oligomer has been developed which is capable of inhibiting expression CC of an interleukin gene. The interleukin gene is preferably an interleukin CC -15 (IL-15) gene. The oligomer can be an oligonucleotide or an CC oligonucleotide analogue. When it is an oligonucleotide analogue it is gelected from protein nucleic acid, morpholino, methylene linkage, CC boronated, and pteridine oligonucleotide analogues. The analogue is CC linked at its 5' end or 3' end to an intercalator. The intercalator is a CC psoralen or acridine derivative. The oligomer is preferably an CC oligonucleotide made of DNA. The oligomer is preferably an CC oligonucleotide derivative, especially a phosphodiester oligonucleotide. The oligonucleotide is at least 5 (preferably 5-50) nucleotides, in CC length. The present sequence represents a specifically claimed CC oligonucleotide of the present invention. The oligoner can be used to alleviate inflammatory polyarthopathy, especially that associated with C rheumatolid arthritis. The oligomer can also be used to alleviate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligomer capable of inhibiting expression of an interleukin gene - is used to alleviate inflammatory poly-arthopathy, especially rheumatoid
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                                                                                                                                                                                                                                                                                                                                                                                                                                      expression interleukin
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AAV37793
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                                                                                                                                       An oligomer has been developed which is capable of inhibiting expression of an interleukin gene. The interleukin gene is preferably an interleukin contains a preferably an interleukin contains a contained by the contained of the preferably an interleukin contained by the contained analogue. The oligomer can be an oligomuclectide analogue it is selected from protein nucleic acid, morpholino, methylene linkage, contained, and pteridine oligonuclectide analogues. The analogue is contained at its 5' end or 3' end to an intercalator. The intercalator is a psoralen or acridine derivative. The oligomer is preferably an continuation of the protein and contained is a phosphodiester, contigonucleotide derivative, especially a phosphodiester oligonucleotide. The oligonucleotide is at least 5 (preferably 5-50) nucleotides, in contigonucleotide of the present sequence represents a specifically claimed coligonucleotide of the present invention. The oligomer can be used to alleviate inflammatory polyarthopathy, especially that associated with the contained arthritis. The oligomer can be used to alleviate inflammatory polyarthopathy, especially that associated with the contained arthritis.
                         Query Match
Best Local Similarity
Matches 17; Conserv
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligomer capable of inhibiting expression of an interleukin gene - is used to alleviate inflammatory poly-arthopathy, especially rheumatoid
                                                                                               Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 20; Page 8; 19pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-OCT-1996;
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                                                                                                                              eosinophilic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interleukin gene; IL-15; inhibitor; oligomer; expression; transcription-inhibiting complex; polypurine-polypyrimidine region; inflammatory poly-arthopathy; rheumatoid arthritis; asthma; ss.
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 5412 AAGAAATAAAAAGCAAGAGAA 5432
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                                 Conservative
                                                                                               BP; 15
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                                                                                                                             inflammation, especially that associated with chronic asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene
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                                                                                             A; 0 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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                                              Score 14.6; DB 1;
Pred. No. 2.3e+03;
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Pred. No. 2.3e+03;
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AAV10466/c
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                                                                                                                                                                                                                                                                 Primers AAV10465-V10492 are used to amplify regions of the human osteosarcoma cell lines MG-63 and SAOS-2 which contain ligands and growth factors and have been designed to cross intron/exon boundaries. The products are used in a process for propagating and maintaining the immature morphology of mammalian haematopoietic cells. The process involves obtaining an enriched population of mammalian haematopoietic cells having the immature morphology of CD34+, HLA-DR+, Thy-1+ and Linand co-culturing this population in the presence of osteoblast cells for between 2 weeks and 8 weeks. The immature cells can be detected by exposing them to an anti-CD34 antibody immunoreactive with the haematopoietic cell antigen CD34, and removing cells that do not immunoreact with the antibody. Such haematopoietic cells can be infused into the blood stream or bone-marrow cavity to treat blood disorders
              30-NOV-1999
                                        AAZ26171;
                                                                AAZ26171 standard; DNA; 21
                                                                                                                                                                                                                                          Sequence 21 BP; 0 A; 11 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maintenance of haematopoietic cells in culture - by co-culturing with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Emerson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human osteosarcoma PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         osteoblast(s).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer;
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            (first entry)
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RESULT 3245
AAZ26812/c
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                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                         graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
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Polymorphism; human; cell viability; loss
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                                                     polymorphic region 1001.
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                                                                                         (first entry)
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inhibitor; cancer; treatment; cell growth;
of heterozygosity; precancerous condition;
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                                                                                                                                                                                                                                                                                                                                                 Score 14.6;
Pred. No. 2.
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This invention describes a novel method for identifying an inhibitor optentially useful for treatment of cancer, where the inhibitor is active con a gene vital for cell growth or viability, and where the gene is consisted to loss of heterozygosity (LOH) in a cancer. The inhibitor is consect to loss of heterozygosity (LOH) in a cancer. The inhibitor is consect the generous condition, by administering to the patient a first allele consected in inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic colls of the patient are heterozygous for the first gene, the inhibitor colls of the patient are heterozygous for the first gene, the inhibitor colls of the patient are heterozygous for the first gene, the inhibitor colls of the genetive on at least one but less than all allelic forms of the gene constitute on at least one but less than all allelic forms of the gene constant somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic constant versus host disease. The method can also be used to remove consignant cells from bone marrow transplants. AAZ25812-Z26825 represent to the method of the invention to the invention constant cells described in the method of the invention constant cells described in the method of the invention constant cells described in the method of the invention constant cells described in the method of the invention constant cells.
    RESULT 3246
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Matches 17
Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
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                                                                                                                                                        Human polymorphic region 587.
                                                                                                                                                                                                       30-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-0041057P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 14 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stanton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.6;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 3247
AAZ26192/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele greenific inhibitor (ASI) targeted to an allele of a first essential gene CC present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor CC is active on at least one but less than all allelic form present in the green tin a population and targets only one allelic form present in the crossler in a population and treatment of LOH disorders, e.g. CC cancers, atherosclerotic plaques, prevention and treatment of LOH disorders, e.g. CC cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometriosis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                              Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                     Human polymorphic region 381.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for identifying an inhibitor
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WO9841648-A2
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                                                                                                                                                                                                                                                                                          AAZ26192
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                                                                                                                                                                                                                                                                                                                                                                                                              5473
                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity
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                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                              TTTTTTTGTAAAAAGATAATT 5493
                                                                                                                                                                                                                                                                                                                                                                              TTTTTTGCAAAAACTAAAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                        (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 2 C; 1 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Đ,
                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                            21
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                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                               cc on a gene vital for cell growth or viability, and where the gene is cc subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is cc used for preventing the development of cancer in a patient having a cc precancerous condition, by administering to the patient a first allele cc precent in cells of the precancerous condition, where the normal somatic cc present in cells of the precancerous condition, where the normal somatic cc lls of the patient are heterozygous for the first gene, the inhibitor cc active on at least one but less than all allelic forms of the gene crownal somatic cells, and the first gene. The products and methods can be cused in the diagnosis, prevention and treatment of LOH disorders, e.g. cc cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic clesions, benign tumours, endometriosis, polycystic kidney disease, and cc malignant cells from bone marrow transplants. AAZ25812-Z26825 represent con the invention sites described in the method of the invention
                                                                         Query Match
Best Local S
                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-MAR-1998;
                                                                                                                               Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                             potentially useful
                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 7; 605pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Housman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (VARI-) VARIAGENICS INC
                            7415
                                                              17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ď
                                                                                Similarity
                    GCAGCAGCAGCAGCAGCA 7435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ledley FD,
                                                                                                                               BP; 2 A;
                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-US005419
                                                                                                                                                                                                                                                                                                                                                                                                                                 for treatment
                                                                            0.2%;
                                                                                                                               7 C; 7 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stanton
                                                              0,
                                                                                Score 14.6;
Pred. No. 2
                                                                                                                               5 T; 0 U; 0 Other;
                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   method for identifying an inniurco.
where the inhibitor is active
                                                                            3e+03
                                                                                                  DB
                                                                                             1.
                                                                                               Length
                                                                  Indels
                                                                                                 21;
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                                                              Gaps
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RESULT 3248
AAZ26714/c

ID AAZ26714;

XX

AC AAZ26714;

XX

AC AAZ26714;

XX

Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH;

KW cell viability; loss of heterozygosity; precancerous condition; ASI;

KW allele specific inhibitor; somatic cell; diagnosis; prevention;

KW allele specific inhibitor; somatic cell; diagnosis; prevention;

KW allele specific plaque; premalignant metaplastic lesion; endometriosis;

KW dysplastic lesion; benign tumour; polycystic kidney disease; transplant;

KW dysplastic lesion; benign tumour; polycystic kidney disease; transplant;

XX

YX

YX

Homo sapiens.

XX

24-SEP-1998;

98WO-US005419.
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AAZ26761/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                     Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viabbility; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ26761 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 15 A; 3 C; 1 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 7; 605pp; English.
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                                                                       19-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ26761;
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(VARI-) VARIAGENICS INC.
                                    20-MAR-1997;
                                                                                                             24-SEP-1998
                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4466
                                                                                                                                                                                                                                                                                                                                                      polymorphic region 950.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                    97US-0041057P
                                                                         98WO-US005419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
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on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is cused for preventing the development of cancer in a patient having a creation condition, by administering to the patient a first allele greeancerous condition, by administering to the patient a first allele concerns to really soft the precancerous condition, where the normal somatic colls of the patient are heterozygous for the first gene, the inhibitor calls of the patient are heterozygous for the first gene, the inhibitor calls of the patient are heterozygous for the first gene, the inhibitor calls contain a population and targets only one allelic forms of the gene contains somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic concerns, benign tumours, endometriosis, polycystic kidney disease, and can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent concerns and contains and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
Sequence 21 BP; 3 A; 12 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is ac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 7; 605pp; English.
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RESULT 3250
AAX17912/c
                                                                                                                                                                        Matches
                                                                                                                                                                                              Query Match
Anti-CMV oligonucleotide #4847.
                        11-MAY-1999
                                                                    AAX17912 standard; DNA;
                                                                                                                                                                                     Local
                                                                                                                                                 2865 AGCAAGGAGGAGGTGGG 2885
                                                                                                                             21
                                                                                                                                                                                   n 0.2%;
Similarity 81.0%;
                                                                                                                                                                         Conservative
                        (first entry)
                                                                     21
                                                                     βP
                                                                                                                                                                        0
                                                                                                                                                                                    Score 14.6; DB 1;
Pred. No. 2.3e+03;
                                                                                                                                                                          Mismatches
                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                               Length 21;
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                                                                                                                                                                         0;
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0,

멼 S

Synthetic.

herpesvirus

'n

modified_base

Location/Qualifiers

WO9845314-A1

linkages"

/*tag= a /note= "contains phosphorothioate internucleotide

(ISIS-) ISIS PHARM INC

09-APR-1997; 07-APR-1998;

97US-00838715 98WO-US006895 Antisense; oligonucleotide; immediate early; DNA polymerase; cytomegalovirus; inhibition; replication; sugar modification; phosphorothioate; infection; retinitis; ss.

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RESULT 3251
AAZ00585
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      glypican-6; glypican-4; glypican-1; glypicatreatment; abnormal; cell growth; cell behatumour formation; sequence tag; STS; MV3a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glypican; GPC1; GPC3; GPC4; GPC5; GPC6; human; glypican-related protein; glypican-6; glypican-4; glypican-1; glypican-3; glypican-5; diagnosis; treatment; abnormal; cell growth; cell behaviour; somatic overgrowth;
This invention describes the isolation of novel human polynucleotides encoding glypican-related proteins, glypican-6 (GPCG) and glypican-4 (GPC4). The invention also describes the polynucleotide and encoded protein sequences of glypican-1 (GPC1), glypican-3 (GPC3) and glypican-5 (GPC5). The products of the invention can be used to diagnose and treat
                                                                                                                                                                                                 e.g.
                                                                                                                                                  Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 methoxyethoxy) sugar modification or phosphorothicate internuclectide linkages. The oligonuclectides are used to inhibit CMV infections (by in vivo or in vitro contact with cells, tissues or body fluids), especially to treat or prevent CMV infections, particularly retinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotides (AAX17861-X17924) are targeted to a nucleic acid (AAX17925-X17948) encoding IE (immediate early) 1 or 2, or DNA polymerase of cytomegalovirus (CNV) and are able to inhibit CNV replication. Optionally the oligonucleotides include at least one 2'-(i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides that target cytomegalovirus nucleic acid particularly including 2-methoxyethoxy sugar modifications, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Draper
                                                                                                                                                                                                                                                                                                                      Veugelers MPD,
                                                                                                                                                                                                                                                                                                                                                                     (VLAA-) VLAAMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9937764-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3024
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                                                                                                                                                                                                                                                                                                                      David
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                                                                        This oligonucletide is complementary to a portion of DNA sequence (see CANV11743) coding for the P30/PHAPI (see AAW84053) of the newly identified CV 3loop HIV receptor. It is used as a sense primer, together with an CV antisense primer (see AAV71752) in a PCR amplification of P30/PHAPI CY reverse-transcribed mRNA. The V3 loop HIV receptor consists of an CV association of 3 proteins, named P95/nucleolin, P40/PHAPI and P30/PHAPI CY (see AAW84052-54). A method for screening molecules that modulate the CV expression of the receptor comprises: cultivating cells transfected with CY a nucleotide sequence encoding P95/nucleolin, P30/PHAPI or P30/PHAPI (c) placed under the control of its own promoter; bringing the cells into CV placed under the control of its own promoter; bringing the cells into CV contact with a test molecule, and quantifying expression of the CV p95/nucleolin, P30/PHAPI or P40/PHAPI e.g. by quantitative PCR using the CV p95/nucleolin, P30/PHAPI or P40/PHAPI e.g. by quantitative pCR using the CV primers provided (see AAV71749-54). Active molecules that have the CV primers provided (see AAV71749-54). Active molecules that have the CV ability to alter and/or prevent the binding of the HIV receptor to the CV of the invention
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Briand J, (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          isolated V3 loop HIV receptor - comprises P95/nucleolin, P40/PHAPII P30/PHAPI proteins, used to develop products for the treatment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
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CENT NAT RECH SCI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 48;
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PI; infection; therapy; PCR; primer;
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Pred. No. 2.3e+03;
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                                                                                      associated bone marrow hypoplasia (e.g. aplastic anemia following themotherapy or bone marrow transplant) may be effectively treated with the antibody compounds as well as disorders such as disseminated intravascular coagulation (DIC), immune thrombocytopenia (HIV-induced and non HIV-induced), chronic idiopathic thrombocytopenia, congenital thrombocytopenia, thrombotic thrombocytopenia and myelodysplasia. They can also be used in e.g. myelotoxic chemotherapy for treatment of solid tumours or leukaemia, myeloablative chemotherapy for autologous or allogeneic bone marrow transplant, myelodysplasia, idiopathic aplastic anemia, congenital thrombocytopenia, and immune thrombocytopenia. The antibodies which bind to the MuSK receptor can be used for improving neuromuscular function in a patient, e.g. in muscular dystrophy. The products can also be used for detection and diagnosis. The antibodies have a longer half-life than the natural ligand for the TPO-R. Sequences
                                                                                                                                                                                                                                                                                                                                                             thrombopoietin receptor (TPO-R). The antibodies which bind the TPO-R can be used in the same way and for the same indications as thrombopoietin (TPO). They can stimulate proliferation, differentiation or growth of megakaryocytes. They may also be able to stimulate megakaryocytes to increase platelet production. They can be used for treating immunological or hematopoietic disorders, especially thrombocytopenia. Thrombocytopenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 megakaryocyte; platelet; immunological; hematopoietic; thrombocytopenia; bone marrow hypoplasia; disseminated intravascular coagulation; anemia; myelodysplasia; myelotoxic chemotherapy; leukaemia; tumour; MuSK; CDR; neuromuscular; muscular dystrophy; complementarity determining region;
                                    AAX32387-X32413 represent DNA fragments encoding the CDR1, CD regions of variable heavy (VH) chains and variable light (VL) antibodies Ab1 to Ab6
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an agonist antibody (Ab) which binds thrombopoietin receptor (TPO-R). The antibodies which bind the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 10; Page 75; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New thrombopoietin receptor agonist antibodies - useful for treating immunological or hematological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Adams CW, Carter PJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC
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                                                        R2, and CDR3
chains of
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or congenital/acquired defects such

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AAZ06694/c
ID AAZ066
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    CC separated by a 1.1 kb intron, and exons 2 and 3 are separated by a 1.6 kb cc intron. The DLX3 gene has been localised to chromosome 17q20-21. The DLX3 grotein has a molecular weight of 32kb. The dlx genes are thought to cf function in the normal cranifacial development and the development of cc mormal dentition. Tricho-Dento-Osseous syndrome (TDO) is an autosomal considered by abnormal hair, teeth and bone. Studies con TDO patients show that they have a mutated form of DLX3 called cc on TDO patients show that they have a mutated form of DLX3 called cc products, this results in a frame shift causing the DLX3delta protein (AAY9927) to be a truncated form of DLX3. The nucleotide and amino acid sequences of both DLX3 and DLX3delta can be used in the development of cc products for enhancing growth, development and repair of osseous structures, particularly for treating bone defects especially in TDO cc sufferers. Addition of DLX3delta proteins and DLX3delta-encoding nucleic caids should be arve to enhance bone thickness and increase bone density at the sites of application. Exogenously added DLX3delta proteins and CC bone/osseous defects secondary to trauma, such as broken bones. Finally, the DLX3delta proteins and nucleic acids should also have utility in the treatment of constituent of defects secondary to certain pathologies, such as contains and afforts and affo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 3254
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primers AAZ06693-Z06694 are used to amplify the coding sequence of DLX7. DLX7 is a member of the distal-less homeobox gene family. DLX7 is used in the isolation of DLX3 (AAZ06690). The DLX3 gene consists of three exons with the homeobox contained in exons 2 and 3. Exons 1 and 2 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated DLX3 proteins and nucleic acids, used to develop products for enhancing growth, development and repair of osseous structures, particularly for treating bone defects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            craniofacial development; Tricho-dento-osseous syndrome; TDO; abnormal hair; teeth; bone; osseous structure repair; bone defect; bone thickness; bone density; broken bone; periodontal disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ06694 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reverse PCR primer DLX7R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             54pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amplification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of DLX7
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syndrome; TDO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 3255
AAX17984
                                                                                    This oligonuclectide was used to generate a variant anti-CD3 antibody

(Ab) heavy chain in a new method for preparing a multispecific Ab

comprising a first polypeptide (PP) and at least 1 extra PP, where: (i)

the first PP comprises a multimerisation domain (MD) forming an interface

positioned to interact with an interface of a MD of the extra PP; and

(1i) the first and extra PPs each have a binding domain, which comprises

ca heavy chain and a light chain, where the variable light chains of the

first and extra PPs comprise a common sequence. The method comprises: (a)

culturing a host cell comprising nucleic acid encoding the first PP and

cextra PP, and the variable light chain, such that the nucleic acid is

expressed; and (b) recovering the multispecific Ab from the culture. The

method prepares heteromultimerric PPs, such as bispecific Abs, bispecific

cundesired formation of the desired heteromultimer relative to the

cundesired heteromultimers and homomultimers. The method allows for the

cundesired heteromultimers and homomultimers. The Abs can be used in

cundiscired heteromultimers and homomultimers. The Abs can be used in

cundiscired heteromultimers in vitro or in vivo diagnosis of various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 17
               Query Match
Best Local 9
   Matches
                                                                                                                                                                                                                                                                                                                                                                                                            Selective preparation of multispecific antibodies - with heteromultimeric heavy chain and common light chain components, useful for, e.g. in vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-MAY-1997;
24-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-070091/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Variant; antibody; heavy chain; light chain; immunoadhesin; immunoassay; diagnosis; cancer; primer; PCR; amplification; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            osteoporosis. The products can also be used for detection and diagnosis
                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer D399S to generate variant humanised anti-CD3 heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
Local Similarity
mes 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4570
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                                                                                                                                                                                                                                                                                                                                                                 1; Page 43; 69pp; English.
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 Conservative
                                                            BP; 3
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97US-0050661P.
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              0.2%;
                                                            7 C; 9
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               Score 14.6; DB 1; Length 21; Pred. No. 2.3e+03;
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Pred. No. 2.3e+03;
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                                                            2 T; 0
Mismatches
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                                                            U; 0 Other;
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Indels
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Gaps
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RESULT 3256
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ID AAX8408
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                                                                                                                                This sequence represents a PCR primer for DNA encoding a turkey coronavirus (TCV) S1 protein. The invention relates to an isolate of TCV obtained from a turkey having spiking mortality of turkeys (SMT) or obtained from an animal exposed to the turkey, which isolate is adapted to cell culture. The TCV and bovine coronavirus isolates are useful for forming vaccines or immunogenic compositions useful for preventing or inhibiting SMT. SMT is a form of poult enteritis and mortality syndrome and causes diarrhoea in turkey poults. The methods are useful in diagnosis and detection of turkey coronavirus associated with SMT. The oligonucleotides are useful as primers for amplification of the coronaviruses, which is useful for detecting the replication or presence of a virus in a sample. N.B. The protein encoded by the amplified sequence is referred to in the specification, but is not given in the
                                                                                               Sequence 21
                                                                                                                        specification
                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                          Turkey coronavirus isolates associated with spiking mortality of turkeys
                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-347478/29
                                                                                                                                                                                                                                                                                                                                                                                              Brown TP,
                                                                                                                                                                                                                                                                                                                                                                                                                     (UYGE-) UNIV GEORGIA RES (BROW/) BROWN T P. (VILL) VILLEGAS P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAY-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9925838-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TCV; S1 protein; turkey enteritis coronavirus; turkey coronavirus; spiking mortality of turkeys; vaccine; immunogenic composition; poult enteritis and mortality syndrome; diarrhoea; turkey poult;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer for TCV S1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX84089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX84089 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-NOV-1997;
10-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Turkey coronavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-AUG-1999
                      2679
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                                               l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                   7; Page 7; 90pp;
                   TGGAGAGGGGAGCCACATATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCGCCTGACCTCCGCACGGG 205
TGGAGAGGGCACCGACTTATC 1
                                                                                                                                                                                                                                                                                                                                                                                           Villegas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer; ss.
                                                                                                BP;
                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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98US-00188979.
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                                                                                             4 A;
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                                                        0.2%;
                                                                                               7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein
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                                                                                                                                                                                                                                                                                                                                                                                              Contreras
                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                               FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                               <u>ი</u>
                                               0;
                                                          Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                INC
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                                               Mismatches
                                                                                           ij,
                                                                                               0
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                                                                                               ₽;
                                                        2.3e+03
                                                                                                0 Other
                                                                      DB 1; Length 21;
                                               Indels
                                               0
                                               Gaps
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RESULT 3257 AAX54710/c ID AAX5471

AAX54710 standard; DNA; 21

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                              S
                                                                                                                                                                     CC directed against at least 2 mRNAs selected from target genes, coding and conon-coding regions of RNAs corresponding to target genes, gene initiation CC codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'CC end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, CC conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55180-271) can be used for the antisense treatment of CC diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and inflammation, including lung CC diseases, pulmonary vasoconstriction, inflammation, including lung CC diseases, pulmonary vasoconstriction, inflammation, respiratory CC distress syndrome, pain, cystic fibrosis, pulmonary hypertension, CC gulmonary vasoconstriction, emphysema, chronic obstructive pulmonary CC disease (COPD), and cancers such as leukemiae, lymphomae, carcinomas e.g. CC colon cancer, breast cancer, lung cancer, pancreatic cancer, metastases, as CC well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer
                                                                    Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; colon cancer; breast cancer; lung cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes antisense oligonucleotides (AAX52869-X55271)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 55; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides used in treatment of, e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-SEP-1997;
09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX54710;
                                                                                                                                              Sequence
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                                  3384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fibronectin antisense oligonucleotide fragment.
21
                                                                    1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer;
                                  CCTCCCCAGCTGCCACCCCC 3404
                                                                                                                                              21
                                                                                                                                              BP; 0
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-0059160P.
98US-00093972.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98WO-US019419
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                                                                                                                                              A,
                                                                                   0.2%;
                                                                                                                                              3 C; 14 G;
                                                                      <u>.</u>
                                                                                       Score 14.6;
Pred. No. 2.
                                                                                                                                              4 T; 0 U;
                                                                      Mismatches
                                                                                         2.3e+03
                                                                                                          DB 1;
                                                                                                                                              0 Other;
                                                                                                        Length 21;
                                                                      Indels
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RESULT 3258

Synthetic

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SXXXXXXXXXXXXX
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AAX09053
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                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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04-AUG-1998;
25-SEP-1998;
                                                                                                                                                                                                                                                                                                                                  used in an assay to screen for compounds that change its binding activity to its receptor. The ligand can also be used for treating HIV infection and AIDS. The present sequence represents a PCR primer used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G protein-coupled receptor protein; APJ; central nervous system; circulation; immune function; gastrointestinal function; reproduction; metabolic function; HIV; infection; AIDS; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a ligand polypeptide for the G protein-coupled receptor, APJ. The APJ ligand can modulate central nervous system function, circulatory function, immune function, gastrointestinal function, metabolic function and reproductive function. An antibody against the APJ ligand can be used in diagnosis. The APJ ligand can be
                        Tumour necrosis factor alpha; TNF-alpha; antisense oligonucleotide; ASO; inhibition; expresssion; treatment; disease; disorder; ss.
                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New ligand polypeptide for modulating central
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hinuma S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-JUL-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX83847;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 11; Page 93; 169pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-405507/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse ligand
                                                               Tumour necrosis
                                                                                        14-JUN-1999
                                                                                                                 AAX09053
                                                                                                                                           AAX09053 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                                                                                   3866 TTCCTCCTACCTCCCGCCCCG 3886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tatemoto K,
                                                                                                                                                                                                                                                                                                                 B₽;
                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptide PCR primer SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry
                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98JP-00032577.
98JP-00220853.
98JP-00271645.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97JP-00353955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-JP005805
                                                                                                                                                                                                                                                                                                                 6 A; 3 C; 11 G; 1 T; 0 U; 0 Other;
                                                              factor alpha antisense oligonucleotide.
                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for the G protein-coupled receptor, APJ, useful nervous system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
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                                                                                                                                                                                                                                                                0
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Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Habata Y,
                                                                                                                                                                                                                                                                                       DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fujii R,
                                                                                                                                                                                                                                                                Indels
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AAA34157/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                 Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotides (ASO) for inhibiting a tumour necrosis factor-
alpha (TNF-alpha) gene in an animal, preferably a human, comprise 12-50
nucleotides, 90% of which are complementary to a region of mRNA
containing a GGA sequence motif. The ASO is used to inhibit expression
of a gene in an animal and for treating the animal when afflicted with a
disease or disorder characterised by the presence of an mRNA from a gene
containing a GGA motif. The ASO are specifically targeted to a GGGA
sequence motif found in mRNA from a gene. A study of known ASO has shown
that at least half of the most efficacious ASO's contain one or more TCCC
motifs. This ASO was designated TJU-2826 and corresponds to a region of
            03-AUG-1998;
                                                                                                                                                                        lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ju G,
                                         03-AUG-1999;
                                                                                                                                                                                                                                                                                             Human adenosine receptor related polynucleotide SEQ ID NO:1846.
                                                                                                                                                                                                                                                                                                                                                                                     AAA34157 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Generation of antisense oligonucleotides - GGGA motif found in mRNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rattus rattus
                                                                                                   WO200009525-A2
                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                             28-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3' untranslated region of the primary transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           566
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                                                                                                                                                            leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTGGGGAAGGGAAGGATCGAA 586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTGAGGGAGGGAAGGAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 32; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 8 A; 1 C; 11 G; 1 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                           (first entry)
             98US-0095212F
                                         99WO-US017712.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97US-0051705P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98WO-US013711
                                                                                                                                                          lymphoma; carcinoma; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
                                                                                                                                                                                                                                                                                                                                                                                      ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       by specifically targeting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                          88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of rat TNF-alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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PCR primer; polymerase chain reaction; amplification; probe; detection; Trachoma chlamydia; Mycobacterium tuberculosis; Hepatitis B virus; Hepatitis C virus; ss.

Oligonucleotide PCR primer #10

04-MAY-2000

(first

entry

CN1232182-A Synthetic.

20-0CT-1999.

13-APR-1998; 13-APR-1998;

98CN-00106616 98CN-00106616 밁 ঠ

21

CCGCCCACAACGCCCACCCCC 1

3384 CCTCCCCCAGCTGCCACCCCC 3404

0

AAZ88382 RESULT 3261

AAZ88382 standard; DNA;

21 ΒP

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Ce e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, crimpeded respiration, respiratory distress syndrome, pain, cystic certificons, pulmonary hypertension, emphysema, chronic obstructive pulmonary hypertension, emphysema, chronic obstructive containmants, and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including contens and prostate cancer. The reduction of the adenosine content of the CONs reduces side effects. The A-containing ONs break down with the crelease of deoxyadenosine which activates adenosine receptors causing connchoconstriction and inflammation. AAA32313 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present content on, which correspond to SEQ ID NO:1 to 2815, and then the last 185 equences are also called SEQ ID NO:1 to 185, but the sequences differ correspond to called SEQ ID NO:1 to 1860 (AAA32323 to AAA3392) are specifically claimed ONs from the present invention N.B. Sequences given in the disclosure of the present invention do not match contained the corresponding SEQ ID NO: sequences given in the sequence
Matches
                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acids involved in bronchoconstriction, diretyres, directions are inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating effects afflict the lungs of a subject. They can be used for treating and the subject of the lungs of a subject.
                                                                                             Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vasoconstruction,
bronchitis, emphy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYEC-) UNIV EAST CAROLINA.
                        Local
17;
                        Similarity
                                                                                             BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        emphysema, respiratory distress syndrome, ischemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 496;
Conservative
                                                                                             Þ
                      81.0%;
                                                                                        3 C; 14 G; 4 T; 0 U; 0 Other;
                                              0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1343pp; English
0
                        Score 14.6;
Pred. No. 2
Mismatches
                   2.3e+03;
                                              DB 1;
                                          Length
Indels
                                              21;
0;
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RESULT 3262
AAA40708/c
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A method has been developed for detecting nucleic acid molecules. The method includes using coating liquid to drop the specific probe onto a detecting membrane (the salt concentration of coating liquid is higher than 2M and the membrane is a high-molecular polymer with average pore size of 1-10 microns); putting one end of the membrane in the liquid containing amplified polymerase chain reaction (PRR) product, and detecting if the amplified PCR product is hybridised with the probe. A reagent box is also disclosed to efficiently and quickly detect Trachoma chlamydia, Mycobacterium tuberculosis, Hepatitis B virus and Hepatitis C virus. AZ288369 to AAZ88383 represent specifically claimed oligonucleotides used in the exemplification of the present invention
                                                                                                                                                                      07-OCT-1998;
28-DEC-1998;
17-MAR-1999;
Nucleic acids encoding mutant CD36 proteins useful for preventing, diagnosing and treating parasitic infections, especially malaria.
                                                                                                                                                                                                                                                                                                                          Human; rat; CD36; SHR; spontaneous hypertensive rat; diagnosis; therapy; screening; polymorphism; variant; detection; mutant; blood; mutation; insulin; glucose metabolism; fatty acid metabolism; catecholamine; malaria; infection; parasite; antiparasitic; antidiabetic; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                           Rat Nos3 primer Nos3R SEQ ID NO:96.
                                                                                                                                                                                                                                                                                                                                                                                                                      15-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA40708 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymerase chain reaction test method and reagent box
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                                        WPI; 2000-303596/26.
                                                                                                                                                                                                                                                                                                     Rattus
                                                                                                                                 (MEDI-)
                                                                                                                                                                                                                         07-OCT-1999;
                                                                                                                                                                                                                                                                           WO200019883-A2
                                                                                                                    (AITM/)
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17; Conserv
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                                                                                          SCOTT J.
STANTON L W.
                                                                                                                                             MEDICAL RES COUNCIL.
                                                                                                                    AITMAN T J.
                                                                                                                                 SCIOS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGGACAGATGAAGAGGAAGTA 1043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGGAAAGGTGAAGGGGCAGTA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.2%;
larity 81.0%;
Conservative
                                                                Scott J,
                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                      98US-00167750.
98US-00221222.
99US-00270542.
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                                                                  Stanton
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Pred. No. 2.3e
0; Mismatches
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Example 1;
Page 111; 167pp;
English
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CC integree the invention describes (B). Parasites such as plasmodium
CC encoding mutant CD36 proteins (B). Parasites such as plasmodium
CC falciparum (the major cause of malaria) are unable to utilise the mutated
CC proteins to gain entry to, and infect cells. The mutant CD36 proteins do
CC cells. The nucleic acids may be used for the recombinant production of
CC mutant CD36 proteins according to standard methodologies. They may be
CC used in this way to prevent and treat parasitic infections that utilise
CC colls (protein to infect cells, such as P. falciparum, the major cause
CC of CD36 protein to infect cells, such as P. falciparum, the major cause
CC of CD36 protein to infect cells, such as P. falciparum, the major cause
CC of CD36 expression and activity or a patient's CD36 DNA may be screened
CC to determine whether there are any mutations present that may confer
CC resistance to parasitic infections. The proteins and nucleic acids may
CC also be used to prevent, diagnose and treat diseases associated with
CC defects in insulin action and/or glucose metabolism and/or fatty acid
CC metabolism and/or catecholamine action in subjects possessing mutations
CC in the CD36 genes. AAA40606 to AAA40759, and AAB02515 to AAB02564,
CC represent nucleotide and amino acid sequences respectively which are used
CC in the examilition of the present invention the exemplification of the present invention invention describes isolated nucleic acid molecules æ

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XXXSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
                    0.2%;
Best Local Similarity 81.0%;
Matches 17; Conservation
                                                    Sequence 21 BP; 6 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
        1673 CTTGTTTCTGCAAATATGCAC 1693
21
CATGITTATGCAGACATGCAC 1
                          0
                                Score 14.6; DB 1
Pred. No. 2.3e+03
                           Mismatches
                                        DB 1;
                                        Length 21;
                           Indels
                           0
                          Gaps
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RESULT 3263
AAZ73907/c
                                                                                             AAZ73907 standard;
Human biallelic marker downstream amplification primer SEQ ID NO:8263.
                               10-SEP-2001
                               (first entry)
                                                                                              DNA;
                                                                                              2
                                                                                              ВP
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Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymotphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; bybridisation; densityping; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer; diagnosis; ss.

Homo sapiens

WO9954500-A2

28-OCT-1999.

21-APR-1999; 99WO-IB000822.

21-APR-1998; 23-NOV-1998; 98US-0082614P 98US-0109732P

(GEST) GENSET

Cohen D, Blumenfeld Z Chumakov

WPI; 2000-013267/01. Novel biallelic markers used ő construct a high density disequilibrium

Claim 8; Page 1992; 2745pp; English.

Ģ,

the human

genome

AAZ65654 to AAZ69578 represent human biallelic markers from the present

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RESULT 3264
AAZ75773
ID AAZ7577
XX AAZ7577
XX Human b
XX Human g
KW Human g
KW diagnos
XX diagnos
XX Homo sa
X
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Best Local Similarity
Matches 17; Conserv
AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-APR-1998;
23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genome;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blumenfeld
                                                                                                                                                                                                                                                                                                                                                                                               human genome.
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                                                                                                                                                                                                                                                                                                                                2390; 2745pp;
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98US-0109732P.
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                                                                                                                                                                                                                                                                                                                                   English
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                construct a high density
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Matches
                                                                        AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ65579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses; they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention
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                                   pharmaceutical agents acting on a disease as well as other N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 322
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                                                                                                                                                                                                                                                                                      Novel biallelic markers
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23-NOV-1998;
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haplotyping; hybridisation; identif
amplification; single nucleotide po
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human genome; biallelic marker; high genomic map; haplotype; phenotype: re
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human biallelic
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                                                                                                                                                                                                                                             9; Page 1623;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              aplotype; phenotype; polymorphic base; genotyping; ybridisation; identification; characterisation; single nucleotide polymorphism; SNP; PCR primer;
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98US-0109732P
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                                                                                                                                                                                                                                                                       genome.
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                                                                                                                                                                                                                                          2745pp;
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                                                                                                                                                                                                                                            English.
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Pred. No. 2.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amplification primer
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                                                                                                                                                                                                                                                                                   a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 Other;
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Best Local S
Matches 17
                                                      Query Match
Best Local S
Matches 17
                                                                                                                                                        AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; haplotyping; hybridiaation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                            map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                             Novel biallelic markers used to construct a high density disequilibrium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-1998;
23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human biallelic marker downstream amplification primer SEQ ID NO:10094.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-2001
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                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                Claim 8;
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                                                                       Similarity
                                                                                                                                              invention
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                      GAAGAAAATGAAACCAGACC 3300
                                                                                                                21
                                                                                                                                                                                                                                                                                                                                                                                Page 2382; 2745pp; English.
GAAGAAACAAGAAACCAATCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blumenfeld M,
                                                                                                                B₽;
                                                         Conservative
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98US-0109732P.
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                                                                                                                A; 5 C; 3
                                                                  0.2%;
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Pred. No. 2.3e+03;
                                                                     Score 14.6;
Pred. No. 2.
                                                                                                                G; 1 T; 0 U; 0 Other;
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                                                         Mismatches
                                                                       2.3e+03
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                                                                                      DB 1;
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                                                                                   Length
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RESULT 3268
AAZ76031
ID AAZ7603
XX
AC AAZ7603
XX

AAZ76031 standard;

DNA;

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ВP

AAZ76031;

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RESULT 3267
AAZ76866/c
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                                          S
                                                                                                                                                                                                                                                  CC invention, which contain a polymorphic base at position 24 of their controlled sequences. AAZ69579 to AAZ7440 represent amplification controlled sequences. AAZ69579 to AAZ77440 represent amplification comprises for the biallelic markers. The bhallelic markers of the invention have a variety of uses they can be used for high density mapping of the comman genome, and in complex association studies and haplotyping studies compositions and methods of the invention can also be useful for the compositions and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from the confidence of the confide
                                                                                         Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     map
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23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 2623; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cohen D, Blumenfeld
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                                                                                                                                                                                       Sequence 21 BP; 9 A; 1 C; 9 G; 2 T; 0 U; 0 Other;
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21
                                                                                           l Similarity
17; Conserv
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                                                                                                                                                                                                                                        invention
CTCCATGTCTCCATTTTCCTC
                                                                                              Conservative
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98US-0109732P.
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                                                                                                                                          Score 14.6;
                                                                                                                      Pred.
                                                                                              Mismatches
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                                                                                                                 2.3e+03
                                                                                                                                          DB 1;
                                                                                                                                        Length 21;
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10-SEP-2001 (first entry)

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RESULT 3269
AAA97479/c
ID AAA97479 standard; DNA; 21
XX
AC AAA97479;
XX
DT 29-JAN-2001 (first entry)
XX
DE Phytolacca americana antif
XX
Phytolacca americana.
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                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                   identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 an 3367, are not actually given a sequence in the Sequence Listing from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genomic map; haplotype; phenotype; polymorphic base; genotyping haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          map
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                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel biallelic markers
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23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                           invention
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                                                                                                                                                                                                                                                                                  TGGTAACATCCCAGCTGGGAC 2408
                                                                                                                                                                                                                                                                                                                                                                            21 BP; 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Blumenfeld
                              americana antifungal protein; Pa-AFP; Virginian pokeweed; production; Escherichia coli; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                          (first entry)
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98US-0109732P
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                                                                           antifungal protein Pa-AFP XbaI 5' PCR
                                                                                                                                                                                                                                                                                                                                                                            5 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          used
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                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English
                                                                                                                                                                                                                                                                                                                               Score 14.6;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                            .3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         high density disequilibrium
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                                                                                                                                                                                                                                                                                                                                            Length 21;
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                                                                             primer.
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06-APR-1999; 24-MAR-2000; 26-OCT-2000

99US-0127958P

2000WO-US008020

Homo sapiens. cancer;

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WO200062736-A2

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RESULT 3270
AAF20279/c
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Best Local S
Matches 17
                                                                                                                                                       Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antisathmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinit; pulmonary hypertension; emphysema; pulmonary transplantation rejection chronic obstructive pulmonary disease; pulmonary infection; bronchitis
                                                                                                                                                                                                                                                                                                                                                 AAF20279;
                                                                                                                                                                                                                                                                                                                                                                        AAF20279 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of Phytolacca americana (Virginian pokeweed), to cDNA encoding it (AAA97476), to the processes and primers used to clone the AFP cDNA from (Yayanian pokeweed seed total mRNA, and to its IPTG-inducible recombinant production in Escherichia coli. Pa-AFP has activity against such fungi as Rhigoctonia solani (black scurf of potato) and may therefore be useful in agriculture. Sequences AAA97477-A97482 represent PCR primers used to introduce restriction sites into cDNA encoding Pa-AFP in the cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recombination antifungus protein gene sequence, engineering bacterium process for preparing its products.
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                                                                                                                                                                                                                                                                                            Human fibronectin polynucleotide fragment #1846.
                                                                                                                                                                                                                                                                                                                        14-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Sim.
hes 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention to an antifungal protein (Pa-AFP; AAB23178) from the
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                                                                                                                                                                                                                                                                                                                      entry)
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Pred. No. 2.
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The present invention describes low adenosine (A) content antisense CC oligonucleotides and compositions (I) comprising them. In the antisense CC oligonucleotides the A is replaced by a 'Universal' or alternative base. CC (I) can have respiratory, bronchodilator, antiinflammatory, analyssic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC The antisense oligonucleotides and (I) can be used to down-regulate the CC expression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and CI cambodines, endogenously produced specific and non-specific enzymes, communoglobulins and antibodies, antibody receptors, cytokines and CC chemokine receptors, adenosine receptors, bradykinin receptors, cytokine and CC chemokine receptors, adenosine receptors, bradykinin receptors, cytokine and CC receptors, CNS and peripheral nervous and non-specific enzymes, cc including proteins and malignancy associated proteins. The CC including proteins and malignancy associated proteins. The CC including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergy(ies) and/or surfactant hypoproduction which are associated with a disease or cc condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary consent and antisense oligonucleotides used in the exemplification of the present human polymucleotide
CC response and malignancy associated with a disease (COPD), pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polymucleotide from pulmonary consent human polymucleotide
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Best Local S
Matches 17
                                                                                                              Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYEC-) UNIV EAST CAROLINA. (NYCE/) NYCE J W.
05-OCT-2000
                                                                                                                                                                                            Single nucleotide
                                                                                                                                                                                                                                                                        AAC71664;
                                                                                                                                                                                                                                                                                                              AAC71664 standard; DNA; 21 BP
                                      WO200058519-A2
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17; Conserv
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llarity 81.0%;
Conservative
                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                            polymorphism
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Pred. No. 2.3e
0; Mismatches
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                                                                                                                                                                                            PCR primer #996
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RESULT 3272
AAC71649/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosts and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                          Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine systemcological system; forensic testing; paternity testing; PC
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Lipshutz RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified it
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(AFFY-)
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                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism
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             WPI; 2000-611722/58
                                   Altshuler D,
Lipshutz RJ,
                                                                          (WHED )
                                                                                                                                       30-MAR-2000; 2000WO-US008440
                                                                                                                                                                  05-OCT-2000.
                                                                                                                                                                                           WO200058519-A2
                                                                                                                                                                                                                     Homo gapiens
                                                                                                                                                                                                                                                                                                                                                                        AAC71649 standard;
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17; Conserv
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                                                                                       WHITEHEAD INST BIOMEDICAL
                                                                           AFFYMETRIX INC.
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Patil N, s
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                                     Cargill M,
Patil N,
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                                      Sklar P;
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klar P;
                                                 Daley GQ,
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Pred. No. 2.3e+03;
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                                                   Ireland
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                                                                                                                                                                                                                                              endocrine system;
ty testing; PCR primer;
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RESULT 3273
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Best Local S
Matches 17
       This sequence represents a target sequence for a microsphere genosensor created using the composition of the invention. The invention relates to a composition which comprises a substrate (other than a fibre optic bundle) with discrete sites on its surface, and microspheres distributed at these sites. The microspheres comprise at least two subpopulations, each with a bioactive agent and an optical signature that identifies the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosts and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                    Composition comprising microspheres with a signature, at discrete sites on substrate, analytes, e.g. nucleic acids.
                                                                                                          Disclosure; Page 29;
                                                                                                                                                                                                                                                                              11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                     bioactive agent; optical signature; interleukin-6; IL-6; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Microsphere genosensor; detect target analytes; potential drug; pollutant; solvent; therapeutic hormone; cytokine; cancer; Alzheimer's disease; toxic bacteria; forensic DNA fingerprinting; det
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                          Walt
                                                                                                                                                                                                                                                                                                         10-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
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17; Conserv
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                                                                                                                                                                                                                        Michael KL;
                                                                                                                                at
9.9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (IL-6) microsphere genosensor target sequence
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Pred. No. 2.3e+03;
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                                                                                                                                                 bioactive agent and optical useful for detecting target
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tic drug; illicit drug;
se; cystic fibrosis;
detect; infection; target;
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Microsphere genosensors

attaching a

Matches Query Match Best Local :

l Similarity 17; Conserv

Conservative

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Mismatches

Indels

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Gaps

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Score 14.6; DB 1, No. 2.3e+03;

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Sequence 21

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AAA94225
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Best Local S
Matches 17
                                       The present sequence is an antisense oligonucleotide directed at the human testosterone-repressed prostate message-2 (TRPM-2, also known as clusterin, sulfated glycoprotein-2 or SGP-2). The sequence was shown to promote the regression of tumours, and oligonucleotides directed at human TRPM-2 can be used in the treatment of tumour cells expressing the TRPM-gene. These include prostate cancer, renal cell cancer and some breast cancer cells. In addition to this, they also increase the chemosensitivity of the cells, meaning that conventional chemotherapy is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (examples include AAA10964-A10968) to the microsphere surface chemistry. A fluorescent dye molecule is attached to the target sequence (examples include AAA10969-A10973) which is in solution. The optically interrogatable signal change occurs with the binding of the target sequences to the microsphere. The composition of the invention is used to detect target analytes and screen large libraries of bioactive agents for those (potential drugs) that bind to the target analytes e.g. pollutants; solvents; therapeutic or illicit drugs; hormones; cytokines; nucleic acids (detecting genes associated with cancer, Alzheimer's disease or cystic fibrosis), antigens; whole cells; to screen foods and water for toxic bacteria; for forensic DNA fingerprinting, for sequencing and for
                              chemosensitivity of the more effective
                                                                                                                                                                                 Example
                                                                                                                                                                                                               the
                                                                                                                                                                                                                             Treating
                                                                                                                                                                                                                                                                                          Gleave
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; testosterone-repressed prostate message-2; TRPM-2; clusterisulfated glycoprotein-2; SGP-2; cancer; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human testosterone-repressed prostate message-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA94225 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detecting
                                                                                                                                                                                                                                                                                                                                                    26-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA94225;
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                                                                                                                                                                                                                                                                                                                        (UYBR-)
                                                                                                                                                                                                            testosterone-repressed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                            prostatic tumors and renal osterone-repressed prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGAAATACCCCAGACCCAAC 3673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mutations;
                                                                                                                                                                               Page 36;
                                                                                                                                                                                                                                                                                        Rennie PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                        BRITISH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                    99US-0121726P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                        COLUMBIA
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                                                                                                                                                                                                                                                                                        Miyake H,
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                                                                                                                                                                               English.
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Pred. No. 2
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                                                                                                                                                                                                                                                                                          Nelson
                                                                                                                                                                                                            messenger-2
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TRPM-2
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                                                                                                                                                                 Query Match
Best Local
                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                 array with a sample, measuring the optical response of each microsphere sensor element, and performing statistical analysis on the measured optical response of the subpopulations. The method is used for detecting target analytes such as nucleic acids, proteins, hormones, lipids, carbohydrates, whole cells, environment pollutants and drugs. The method is inexpensive when compared to conventional methods. The array once loaded can be decoded or used after testing. The present sequence represents a probe used to detect the presenc of a disease- associated gene fragment in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method for detecting a target analyte in a sample using a self-encoding sensor array comprising a population of microspheres on discrete locations on the surface of a substrate. The array has several subpopulations of microspheres, each of which provides a characteristic optical response signature when illuminated by excitation light energy in the presence of a reference analyte, which may in some cases be the target analyte. The method involves contacting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Method for detecting target analyte in sample involves, providing array with several sub population of sensor elements, measuring optical response of each sensor element and statistically analyzing optical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Analyte detection; microsphere array; self-encoding sensor array; nucleic acid detection; biochip; disease-associated gene; probe; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA97685 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-656240/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 19; Fig 22; 99pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-2000; 2000WO-US009183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200060332-A2
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                                                                                                                                                          Local
                                                             3653 ААСАЛАТАССССАСАСССААС 3673
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                                                                                                                              17; Conserv
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                                                                                                                                                                                                                                                             21
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   AATAACCACCCCTGACCCAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dickinson TA;
                                                                                                                                                                                                                                                             BP; 2 A; 1 C; 10
                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-00287573
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                                                                                                                                                             81.0%;
                                                                                                                                                                                                0.2%;
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                                                                                                                              0,
                                                                                                                           Score 14.6; DB 1;
Pred. No. 2.3e+03;
0; Mismatches 4;
                                                                                                                                                                                                                                                             G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                         Length 21;
                                                                                                                                  Indels
                                                                                                                              0
                                                                                                                              Gaps
                                                                                                                              0,
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RESULT 3276 AAA97679 ID AAA9767

AAA97679 standard; DNA; 21 BP

15-FEB-2001 AAA97698;

(first entry)

AAA97698 standard; DNA; 21 BP

Interleukin-6 (IL-6) array probe #18.

Analyte detection; microsphere array; self-encoding sensor array; nucleic acid detection; biochip; disease-associated gene; probe; ss.

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RESULT 3277
                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method for detecting a target analyte in a cample using a self-encoding sensor array comprising a population of microspheres on discrete locations on the surface of a substrate. The carray has several subpopulations of microspheres, each of which provides a characteristic optical response signature when illuminated by excitation light energy in the presence of a reference analyte, which may in some cases be the target analyte. The method involves contacting the carray with a sample, measuring the optical response of each microsphere contacting the sensor element, and performing statistical analysis on the measured optical response of the subpopulations. The method is used for detecting target analytes such as nucleic acids, proteins, hormones, lipids, carbohydrates, whole cells, environment pollutants and drugs. The method is inexpensive when compared to conventional methods. The array once loaded can be decoded or used after testing. The present sequence conversion a target sequence which can be detected using the method of
                                                                                                                                                                                                                Best Local Similarity Matches 17; Conservat
                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Method for detecting target analyte in sample involves, providing array with several sub population of sensor elements, measuring optical response of each sensor element and statistically analyzing optical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 8 A; 10 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Walt DR, Dickinson TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-OCT-2000.
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                                                                                                                                                     3653 AAGAAATACCCCAGACCCAAC 3673
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                                                                                              AATAACCACCCCTGACCCAAC 21
                                                                                                                                                                                                                0.2%;
milarity 81.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 39; 99pp; English.
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                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                       Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                       .3e+03
                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                          Indels
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AAA65238
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       array with a sample, measuring the optical response of each microsphere sensor element, and performing statistical analysis on the measured optical response of the subpopulations. The method is used for detecting target analytes such as nucleic acids, proteins, hormones, lipids, carbohydrates, whole cells, environment pollutants and drugs. The method is inexpensive when compared to conventional methods. The array once loaded can be decoded or used after testing. The present sequence represents a probe used to detect the present of a disease- associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sample using a self-encoding sensor array comprising a population of microspheres on discrete locations on the surface of a substrate. The array has several subpopulations of microspheres, each of which provides a characteristic optical response signature when illuminated by excitation light energy in the presence of a reference analyte, which may in some cases be the target analyte. The method involves contacting the
                                                                                                                                                                     Meloidogyne
                                                                                                                                                                                                                                                         AAA65238 standard; DNA;
                                          13-JUL-2000
                                                                                                Meloidogyne
                                                                                                                                          Species-specific oligonucleotide; crop
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method for detecting a target analyte in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method for detecting target analyte in sample involves, providing array with several sub population of sensor elements, measuring optical response of each sensor element and statistically analyzing optical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Walt DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-APR-2000; 2000WO-US009183
            28-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                            3653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fragment in an exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                          AAGAAATACCCCAGACCCAAC 3673
                                                                                                                                                                                                                                                                                                                               AATAACCACCCCTGACCCAAC 21
                                                                                                                              nematode;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dickinson TA;
                                                                                                                                                                     chitwoodi species-specific oligonucleotide #1.
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                                                                                                 chitwoodi
                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                 (first entry)
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             99WO-NL000812
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                                                                                                                              PCR
                                                                                                                                                                                                                                                                                                                                                                                                    0.2%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                10 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                            primer; ss
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Pred. No. 2.
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                                                                                                                                          crop damage;
                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                       Indels
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Best Local :
                                                                                                                                                                                                                                                     PCR primer; prostate cancer cell line; androgen independent; CL-1; LNCaP cell line; beta-actin; Prostate-specific antigen; Prostate specific membrane antigen; Basic fibroblast growth factor; Vascular endothelial cell growth factor; Interleukin-6; Transforming Growth Factor-betal; Transforming Growth Factor-beta-R; Epidermal growth factor receptor; PSA; R; PSAM; IL-8; VEGF; bFGF; IL-6; TGF-betal; TGF-beta2; TGF-beta-R; EGF-R; BCL-2; E-cadherin; p53; PTEN; Caveolin; c-myc; HER-2/neu; p27; androgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     knot nematode Meloidogyne chitwoodi. This is a crop parasite which can cause damage to crops such as potatoes, beets, black salsifies and carrots, the damage being so great that in Europe the organism has been given a quarantine status. The oligonucleotide was identified using random amplified polymorphic DNA and subjecting it to a series of sequence can be used in tests to determine both the presence and species of Meloidogyne parasites, which is useful for seed export and also in the
                                                             Belldegrun
                                                                                                                28-JAN-1999;
                                                                                                                                         28-JAN-2000; 2000WO-US002223.
                                                                                                                                                                   03-AUG-2000.
                                                                                                                                                                                           WO200044879-A1
                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                              Androgen
                                                                                                                                                                                                                                                                                                                                                                                                         08-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA76069;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA76069 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a species-specific oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 20; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel DNA oligonucleotide specific for Meloidogyne detect specific Meloidogyne species in a sample.
                                                                                        (REGC ) UNIV CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                              beta-actin PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-DEC-1998;
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                                      2000-499329/44.
                                                                                                                                                                                                                     sapiens.
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for resistance to the parasite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGGAGGCCAGCTGGAGGAAGA 4751
                                                                                                                                                                                                                                            receptor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGGAGAGCAGCAGGAGAAAGA
                                                             AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 9 A; 2 C; 9 G; 1 T; 0 U;
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Pred. No. 2
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Androgen independent, aggressively tumorigenic prostate cancer cell lines designated CL-1 and CL-2, useful as tools for studying the cellular and

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RESULT 3280
AAF96869/c
ID AAF9686
XX AAF9686
XX AAF9686
XX U6-JUN-
XX U6-JUN-
XX Human;
KW polymox
KW myocard
XX myocard
XX myolymox
XX Humona
XX W20111
FT Variati
AX W020011
XX M020011
XX M020011
XX M020011
XX M020011
XX Lander
XX Lander
PR Lander
VX MPI; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity Matches 17; Conserv
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Best Local :
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26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
     WPI; 2001-226749/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human gene single nucleotide polymorphism #1630.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dependent state to the metastatic androgen-independent state
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                                                                                                                                                                                                                                                                                    07-SEP-2000;
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                                                    ES,
                                                                                                      WHITEHEAD I
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ilarity 81.0%;
Conservative
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                                                    Gargill M,
                                                                                                                                                                             99US-0153357P.
2000US-0220947P.
2000US-0225724P.
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                                                                                                      INST BIOMEDICAL M PHARM INC.
                                                       Ireland JS,
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Pred. No. 2
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                                                    Bolk S,
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AAF97449/c
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Best Local (
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                                                                                                                                        10-SEP-1999;
26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the disease, stroke, peripheral vascular diseases, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes an
                                                                                                                                                                                                                                                                                                                                                                  Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembol pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF97449 standard; DNA; 21 BP
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                                                                                                                                                                                                                       15-MAR-2001.
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                                                                                                    WHITEHEAD INST BIOMEDICAL RES MILLENNIUM PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  single
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; 2000US-0220947P.
; 2000US-0225724P.
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                                                                          Bolk S,
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                                                                          Daley GQ,
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                                                                            Mccarthy
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etes and
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Nucleic acids comprising single nucleotide polymorphisms, us applications such as forensics, paternity testing, medicine, analysis and phenotype correlations to diseases such as diab

diabetes

genetic etes and

useful in

WPI; 2001-226749/23.

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RESULT 3282
AAF9724
AAF9724
XX AAF9724
XX AAF9724
XX O6-JUN-
XX Human;
KW polymox
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Best Local S
Matches 17
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                                                                                Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                              10-SEP-1999;
26-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pulmonary embolism; paternity test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
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Example;

Page 185;

242pp;

English

The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided

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26-JUL-2000;
16-AUG-2000;
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Variation
                                                                                               applications such as for analysis and phenotype
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17; Conserv
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                                                                                                acids comprising single nucleotide polymorphisms, useful tions such as forensics, paternity testing, medicine, gene and phenotype correlations to diseases such as diabetes
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also
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                                                                                                                                                                                                                                                                                                                               Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
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                                                                                                                                                                                                                                                                                                                                    Nucleic acids comprising single nucleotide polymorphisms, usuapplications such as forensics, paternity testing, medicine, analysis and phenotype correlations to diseases such as diabatherosclerosis.
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16-AUG-2000;
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype

correlations to diseases. the human gene SNPS shown

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                                                                                                              The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
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26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease, stroke, peripheral vascular diseases, venous thromboembolism (pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                     The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular disease, venous thrombosmbolism and disease, stroke, peripheral vascular disease, venous thrombosmbolism and
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                          Sequence
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26-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human gene single nucleotide polymorphism
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-SEP-2000; 2000WO-US024503
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                       Similarity
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                                                                                          BP;
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2000US-0220947P.
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                   0.2%;
                                                                                       5 C; 2 G;
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                                                                                                                                The present sequence in the specification
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                     Score 14.6;
Pred. No. 2
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                     2.3e+03
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                                                                                          0 Other;
                                          DB 1;
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Matches 17
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26-JUL-2000;
16-AUG-2000;
                                                                                                                              The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                        Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
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21
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17; Conserv
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 CCTTCAGGTACACCACTGGGG
                         CCATGAGGTACACCCCTGTGG 739
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2000US-0220947P.
2000US-0225724P.
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RESULT 3289

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AAH62434;

RESULT 3290 AAH62434

AAH62434 standard; DNA;

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Best Local Similarity
Matches 17; Conserv
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26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                             The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and disease, stroke, peripheral vascular diseases, venous thromboembolism and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; variant thrombospondin 1; variant thrombospondin 4; polymorphism; vascular disease; coronary artery disease; for myocardial infarction; atherosclerosis; stroke; venous through pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                          Nucleic acids comprising single nucleotide polymorphisms, useful applications such as forensics, paternity testing, medicine, gene analysis and phenotype correlations to diseases such as diabetes
                                                                                                                                           pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of
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                                                                                                   Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                             the human gene SNPS shown
                                                                                                                                                                                                                                                                                       Example; Page 57; 242pp; English.
                                                                                                                                                                                                                                                                                                                  atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                    (WHED )
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                        603
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                   CAAGTGGCTAGCCATTGTGAG
CAAGTGGCTGGCCCTGGAGAG
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; 2000US-0220947P.
; 2000US-0225724P.
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                                                  Conservative
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                                                              0.2%;
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Pred. No. 2.3e
0; Mismatches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                           DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing an nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing, medicine and containing oligonucleotides are useful.
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heart (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
Homo sapiens
                       Single nucleotide polymorphism; SNP; human; cancer; heart disease; paternity testing; forensic science;
                                                               SMRT polymorphism containing DNA fragment #176.
                                                                                                                      AAH62275;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cholinergic receptor polymorphism containing DNA fragment #335.
                                                                                            12-SEP-2001
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                                                                                                                                               standard;
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                                                                       Key
Variation
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Variation
                                                                                                               Homo sapiens
                                                                                                                                         heart disease;
                                                                                                                                                     Single nucleotide polymorphism; SNP; human; cancer;
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                                                                                                                                                                                                                                                          AAH62422 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     applications such as phenotype correlation, forensics, paternity testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 43; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
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                                                                                                                                       paternity testing;
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                                                                      Location/Qualifiers replace(11,T)
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RESULT 3293
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Secreted protein; human; PRO protein; neoplastic cell growth; tumour; proliferation; leukaemia; lymphoid malignancy; inflammatory disorder; angiogenic disorder; immunologic disorder; PRO7476; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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17; Conser
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99US - 0138385P
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                                                     This sequence represents a PCR primer used to isolate DNA encoding human CC PRO7476 protein of the invention. The PRO proteins are secreted proteins. Antagonists or antibodies of PRO polypeptides are useful for diagnosing CC and treating tumours are associated with gene amplification, neoplastic CC cell growth and proliferation in mammals, and those conditions (CC characterised by overexpression and/or activation of the amplified genes. CC Such conditions include benign or malignant tumours (e.g. renal, liver, CK kidney, bladder, breast, gastric, ovarian, colorectal, prostate, CC kidney, bladder, breast, gastric, ovarian, colorectal, prostate, CC glioblastomas and various head and neck tumours); leukaemias and lymphoid cmalignancies, neuronal, glial, astrocytal, hypothalamic, and other CC glandular, macrophageal, epithelial, stromal and blastococlic disorders; and inflammatory, angiogenic and immunologic disorders. These may further CC be used to qualitatively or quantitatively detect the expression of corocteins encoded by the amplified genes, and in tumour diagnostics or preparation of a medicament in the treatment of a condition, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumors are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibodies against PRO polypeptides, useful for diagnosing and treating tumors are associated with gene amplification, neoplastic cell growth a
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                                       responsive to the PRO polypeptide, its antagonist
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BP; 3
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  C; 5 G; 4 T; 0 U;
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  0 Other;
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                                            or anti-PRO antibody
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밁 Ś Best Matches Query Match Local 5166 21 CAGGGCCAGTGGGAGCTGCAT 1 l Similarity 17; Conserv CTGGGACAGTGGGCTCTGCAT 5186 Conservative 0.2%; 0, Score Pred. Mismatches 14.6; No. 2. DB 1; Length 21; Indels 0, Gaps 0

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Human interleukin-6 (IL-6) PCR primer,
                        05-JUN-2001
                                               AAF76187;
                                                                        AAF76187 standard; DNA;
                       (first entry)
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                                                                        ВP
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 NO:53.
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IL-7; IL-6; SCF; LIF;

AAF76187/C
ID AAF76188
XX AAF7618
XX AAF7618
XX DT O5-JUNXX Transgg
XW Transgg
XW Transgg
XW Stem Ce
XW Stem Ce
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XW Human i
XX Huma Transgenic mouse; immunodeficient; tissue recipient; lymphocyte deficient; human cytokine; interleukin; IL-7; IL-6; stem cell factor; leukaemia inhibitory factor; GM-CSF; M-CSF; granulocyte macrophage-colony stimulating factor; macrophage-colony stimulating factor; major histocompatibility compation; human monoclonal antibody generation; haematopoietic cell develuman immune system animal model; PCR primer; 88. 30-AUG-2000; 2000WO-US023971. 31-AUG-1999; 08-MAR-2001. WO200115521-A1 Homo sapiens. 99US-0151688P cell development;

(GEMV) GENENCOR INT INC

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC disrupted, causing it to lack mature B and T cells. In particular, both alleles of the recombination activation gene-2 (RAG-2) gene are CC disrupted, which in turn prevents VDJ recombination. The mouse also CC comprises donor (e.g., human) specific transgenes encoding the cytokines (LIF), granulocyte macrophage-colony stimulating factor (SCF), leukaemia inhibitory factor (LIF), granulocyte macrophage-colony stimulating factor (SCF), leukaemia inhibitory factor (LIF), granulocyte macrophage-colony stimulating factor (GM-CSF), and IL-6, which enable it to support the growth of transplanted donor cells. In another embodiment CC finitocompatibility complex (MHC) class II DR3 molecule, where the CC histocompatibility complex (MHC) class II DR3 molecule, where the CC transgene has naturally linked DRab and DQab alleles. The transgenic CC donor, e.g., non-human, macromolecules; to determine the effect compounds have on a human immune system; to generate fully human polyclonal or CC monoclonal antibodies to specific antigens; to determine whether compounds in the investigate the human cell mediated response in a human cCC pathogens and other immunomodulatory compounds; and to determine the calls. The transgenic mouse supports the functional cCC which produce functionally impaired haematopoietic cells or are CCC monoclonal of calls. The transgenic mouse supports the functional cCC which produce functionally impaired haematopoietic cells or are couse provides in the periphery that exhibit MHC-restricted antigen- specific responses. CCC more closely resembles normal ontogeny, as they possess CD4+ T cells in the periphery that exhibit MHC-restricted antigen- specific mice used in the development of human cytokine-expressing transgenic mice
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Best Local S
Matches 17
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                                                                                                        Vascular endothelial growth factor; VEGF; antisense; angiogenesis; cell proliferation; Kaposi's sarcoma; cancer; melanoma; cytostatic antisense therapy; PIGF; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                   P1GF gene
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17; Conserv
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Pred. No. 2.3e+03;
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; sporiasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention provides a composition comprising one or more antisense oligonucleotides directed against vascular endothelial growth factor (VEGF) where the antisense oligonucleotides inhibits proliferation of cells exhibiting autocrine VEGF activity at an IC 5 0 concentration of between 0.5-2.5 micro Ma. The antisense oligonucleotides may be directed against VEGF for inhibiting cancer cell proliferation and angiogenesis. Preferably the oligonucleotide AAH23032 (a modified version of AAH22984) is used and may be utilized to treat Kaposi's sarcoma, ovarian cancer, prostrate cancer, pancreatic cancer or melanoma. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JAN-2001; 2001WO-US000019
                                           26-OCT-1999;
                                                                       26-OCT-2000; 2000WO-US029500
                                                                                                                                                                        Synthetic.
                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                    basal cell carcinoma; seborrheic wart; vitreoretinopathy;
                                                                                                                                                                                                                                                                                                                                                                                               PDGF B hairpin/hammerhead ribozyme recognition site SEQ ID NO:4501.
                                                                                                                                                                                                                                                                                                                                                                                                                                  10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            077/c
AAH62077 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 5 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prostrate cancer, pancreatic carepresents a PIGF gene fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense oligonucleotides useful for inhibiting vascular endothelial growth factor expression, angiogenesis and for treating cancer, e.g., Kaposi's sarcoma, ovarian cancer and prostrate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-451898/48
                                                                                                          03-MAY-2001.
                                                                                                                                        WO200130362-A2
                                                                                                                                                                                                                     sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH62077;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Fig 17b; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gill PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GILL/) GILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5175 TGGGCTCTGCATGTTCTCCAC 5195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGGGCTGAACATGTGCTCCAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Masood R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                             99US-0161532P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                                                                                                     cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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(IMMU-) IMMUSOL INC

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RESULT 3297
AAH24644/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present inventor.

Skin or eye disease and scarring. The metnou inventor involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which cleaves RNA encoding a cytokine involved in ribozyme (I) which control ribozyme (I) which control ribozyme (I) which cytokine (I) which control ribozyme (I) which cytokine (I) w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Robbins JM,
                              protein, useful fish diseases.
                                                                Novel recombinant Rainbow trout galectin protein and gene protein, useful for studying fish immunity mechanism, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                     01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oncorhynchus mykiss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rainbow trout; galectin; cytostatic; antiinflammatory; immunity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rainbow trout galectin cDNA primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH24644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2001069976-A
                                                                                                                                                                                                                                                        (NORQ ) NORINSUISANSHO YOSHOKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell movement; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1321 GCTCCAGACAGACAGGAGGAG 1341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease; cell adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCAGCAGAGAGACAGGAAGAG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                    99JP-00247204
                                                                                                                                                                                                                                                                                                                                                                                                     99JP-00247204.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibition; inflammation; cancer; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                           KENKYU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
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                                                                diagnosis
                                                                                                          encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>;</u>
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Best Local S
Matches 17
The present sequence was used in reverse transcriptase polymerase chain reaction (RT-PCR) analysis of human prostate cancer cells. The invention relates to an isolated polymucleotide comprising a prostate-specific chimeric enhancer (PSE) sequence and a proximal promoter sequence operably linked to a nucleic acid segment that encodes a heterologous polypeptide. The PSE contains an ARE and specifically activates transcription of the nucleic acid segment in a mammalian prostate cell. The construct is useful for the treatment of a prostate disorder or a metastasised prostate cancer, such as hyperplasia or hyperproliferation of prostate cells. It is also useful for directing the tissue-specific expression of a heterologous polypeptide in a human prostate cell. The construct may be administered by injection, infection, transformation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a recombinant protein having a 341 amino acid sequence fully defined in the specification, or its mutant in which at least one amino acid is deleted, substituted or added, but which retains the galectin activity. The protein is useful for studying mechanisms of fish immunity and for diagnosing fish diseases. The galectin proteins are involved in inhibition of cell adhesion, inflammation, metastasis of tumour cells, T cell movement, and apoptosis. The present sequence is a primer which was used in the production of a probe used to analyse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; androgen response element; ARE; cytostatic; gene thera prostate-specific chimeric enhancer; transcriptional regulatitargeted gene expression; prostate cancer; prostate disorder; prostate-specific antigen; PSA; beta-actin; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 4 A; 10 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                     New polynucleotide, useful for treating prostatic cancer, comprises prostate specific chimeric enhancer and proximal promoter sequence operably linked to nucleic acid encoding heterologous polypeptide.
                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF82667 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression of the rainbow trout galectin gene
                                                                                                                                                                                                                                                                                                                                                                              Wu L,
                                                                                                                                                                                                                                                                                                                                                                                                                   (REGC ) UNIV CALIFORNIA SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-OCT-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-OCT-2000; 2000WO-US028444.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200127256-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF82667;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5552 GCAGATGGAGAAGTGGTGTTG
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                                                                                                                                                                                                                                                                                                                                                                                Carey MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                    Page 73; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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99US-0159730P.
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al regulation;
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RESULT 3299
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The invention relates to Detecting a collagen gene alteration associated with a pathological condition in a human subject by obtaining from the subject a sample nucleic acid containing a portion of at least 15 consecutive nucleotides of the segment of the COLIA1 gene extending in the 5' to 3' direction from 78 nucleotides of intron 27 located adjacent exon 28 through the 3' end of intron 51, where the portion contains an contronic nucleotide and a first and second site, determining the sequence of the portion and comparing the sequence of the portion with the corresponding consensus sequence of the COLIA1 gene where a difference between the sequence of the portion and the consensus sequence indicates the presence of the collagen alteration in the subject. The method is used for deterting abnormalities in a COLI or COL9 gene is useful for determining whether a subject is afflicted with pathological conditions associated with an altered collagen gene such as osteoporosis, multiple epiphyseal dysplasia, osteogenesis imperfecta, shortness of stature and low bone density. Identification of an abnormality in a collagen gene is also useful for designing a therapeutic nucleotide or gene therapy agent which can be administered to the subject to correct or alleviate the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         multiple epiphyseal stature and low bon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting collagen gene alteration, useful for diagnosing osteoporosis, multiple epiphyseal dysplasia, osteogenesis imperfecta, shortness of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ala-Kokko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYAL-)
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13-MAR-1994;
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Larson A, Pack M, Colige A, Early
Annunen S, Pihlajamaa T, Vuoristo
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94US-00212322.
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         The invention relates to an isolated Fanconi anaemia protein complex (C (FANCD2) polypeptide. The FANCD2 protein comprises a sequence of 1472 c amino acids fully defined in the specification, its 90% identical CC sequence, a sequence encoded by a polynucleotide that is at least 90% cidentical to sequence siven in specification such as a 5127 base pair CC sequence, or a fragment which is at least 50 amino acids in length. The CC FANCD2 protein is useful for treating an FA pathway defect in a cell CC target or for treating a patient with a defective FANCD2 gene. The FANCD2 CC gene is useful for making a recombinant expression vector. The FANCD2 CC grotein and its gene are useful as a novel target for therapeutic CC associated with DNA repair and cell cycle abnormalities such as Fanconi CC anaemia, Bloom's syndrome, Cockayne's syndrome, Hereditary non-polyposis CC colon cancer, ataxia telangiactasia and Xeroderma pigmentosum. The FANCD2 gene is useful in producing probes and primers for screening patients in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                        Claim 8; Page 56; 103pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         preparing an experimental mouse model for use in screening new therapeutics for treating conditions involving defective DNA repair, and in gene therapy methods. A recombinant vector containing the RANCD2 gene of the invention is useful in gene therapy. This polynucleotide sequence represents a PCR primer for amplifying a FANCD exon relating to the
 The invention relates to an isolated nucleic acid from a human gene encoding aminopeocidase P (XPNEP2), bradykinin receptor B1 (BDKKB1), tachykinin receptor B1 (TACR1), C1 esterase inhibitor (C1NH), kallikrein (KLK1), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme
                                                                                                             New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
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                                                                                                                                                                                                                                                                    SQUIBB
                                                                    977pp;
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                                                                                                                                                                                                                Zerba
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Pred. No. 2.3e
0; Mismatches
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                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA sequence #54.
                                                                                                                                                                                                                쯦
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                                                                                                                                                                                                                Ma-Edmonds
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WO200261131-A2

Chronic

obstructive pulmonary

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Best Local (
                                                                                                                                                                                                                                                                                                 Aminopeptidase P; XPNBP2; bradykinin receptor B1; ds; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; C1NH; kallikrein KLK1; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACE2; protease inhibitor 4; PI4; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; angioedaema; aneurysm; embolism; thrombosis; coronary arrery disease, angioedaema;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     artery disease, arteriosclerosis and/or atherosclerosis, and artery disease, arteriosclerosis and/or atherosclerosis, and artery hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory hypersensitivity reactions, sepsis, autoimmune diseases, inflammatory hypersensitivity reactions, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease (COPD) and enterocolitis (many other diseases and disorders are listed in the specification). The polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21
                                                                                                                                                                                                                     arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; autoimmune disease; inflammatory arthritis; cancer; wound; viral infection; bacterial infection; fungal infection; COPD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABS60171;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS60171 standard; DNA;
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81.0%;
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Pred. No. 2
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03-DEC-2001; 2001WO-US047235

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CC 1 (KLK1), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme CC 2 (ACE2) or protease inhibitor 4 (PI4), comprising at least one CC polymorphic position. Also included are (1) a probe that hybridises to a CC polymorphic position as provided in the detailed summary of single CC nucleotide polymorphisms comprising additional 5 and 3 flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more individuals and determining the encoding a protein selected from the group above; (3) constructing (M2) the plotypes using the genes comprising grouphic positions in a gene encoding a protein selected from the group above; (3) constructing (M2) (4) identifying (M3) an individual at risk of developing a disorder comprises one or more polymorphic positions within a gene encoding a minimistration of an ACE inhibitor and/or vasopeptidase inhibitor comprises one or more polymorphic positions within a gene encoding a individual solution and protein selected from the group above; and (6) genotyping (M4) an individual at protein selected from the group above; and (6) genotyping (M4) an individual approtein selected from the group above; and (6) genotyping (M4) an individual approtein selected from the group above; and (6) genotyping (M4) an individual approtein selected from the group above; and (7) genes which consist one position with a known data set. The genes (M1, M2, M3 and M4) and compositions are useful for detecting, diagnosing, treating, are compositions are useful for detecting, diagnosing, treating, the provided and compositions are useful for detecting, diagnosing, treating, and compositions are useful for detecting, diagnosing, treating, and mysertension, heart failure, myocardial infarction, ventricular consistivity reactions, sepsis, autoimmune diseases, inflammatory hypertensis, inflammatory reactions. sepsis, autoimmune diseases, inflammatory
                                                                                                                                          Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                polynucleotides are also useful for chromosome identification. Antibodie, against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is included in the sequence listing but is not referred to anywhere else in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypersensitivity reactions, sepsis, autoimmune diseases, inflammatc arthritis, cancer, wounds, viral, bacterial or fungal infection, Chobstructive pulmonary disease (COPD) and enterocolitis (many other diseases and disorders are listed in the specification). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid with at least one polymorphic position, us for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tsuchihashi Z, Hui
Swanson BN, Powell
                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated nucleic acid from a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 709; 977pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-2000;
23-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tachykinin receptor B1 (TACR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                encoding aminopeptidase
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                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                    21 BP; 1
TCTCTACTTCCCTCCCTTTGC
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                                                                                                                                          Conservative
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2001US-0263678P
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                                                                                                                                                                        0.2%;
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                                                                                                                                                                                                                                                                                    C; 1
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CR1), C1 esterase inhibitor (C1NH),
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                                                                                                                                                                        Score 14.6;
Pred. No. 2.
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21
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                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                T; 0 U;
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                                                                                                                                                                           3e+03
                                                                                                                                                                                                             B
                                                                                                                                                                                                                                                                                    0 Other;
                                                                                                                                                                                                      Length 21;
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                                                                                                                                      <u>.</u>
                                                                                                                                      Gaps
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RESULT 3303 ABS68528/c ID ABS6852

ABS68528

standard; DNA;

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06-SEP-2001; 2001WO-US028014

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a DNA sequence encoding clock protein BMAL2 (brain-muscle-Arnt-like protein 2). The gene and protein are applicable in diagnosis of and development of drugs for insomnia and other sleeping disorders e.g. non-24-hour sleep, sleep-phase forward or retreat syndrome and time-zone variation syndrome. ABS68501-ABS68552 represent BMAL2 coding sequences and PCR primers of the invention
                                                                                                     Human; oligonucleotide label-domain; CMV; cytomegalovirus; Epstein-Barr virus; lambda-immunoglobulin light chain; hap kappa-immunoglobulin light chain; repetitive Alu sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clock gene Baml2 and expressed clock protein BMAL2 important in clock oscillation mechanism and relating to circadian rhythm, used in diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; clock protein BMAL2; brain-muscle-Arnt-like sleeping disorder; non-24-hour sleep; sleep-phase f
                                                                                                                                                     In-situ analysis synthetic
                                                                                                                                                                                                                            ABK70498 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified.
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                                                                                                                                                                                                                                                                                                  4996 CCAGCTGAAGAACAGAATGGA 5016
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                                                                                                                                                                                                                                                                                                                                     1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  developing drugs for insomnia and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Page 35; 187pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorder; non-24-hour sleep;
syndrome; time-zone variation
                                                                                                                                                                                                                                                                                                                                                                                   21
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                                                                                            early
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                                                                                             RNA;
                                                                                                                                                                                                                                                                                                                                               0.2%;
                                                                                                                                                                                                                                                                                                                                                                                   6 C; 4 G;
                                                                                                                                                    probe
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                                                                                                                                                                                                                                                                                                                                               Score 14.6; DB 1;
Pred. No. 2.3e+03;
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                                                                                                                                                                                                                                                                                                                                      Mismatches
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syndrome; P(
                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein 2)-related
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ABK87131/c
ID ABK8713
XX ABK8713
XX O7-OCT-
XX O7-OCT-
XX Human;
KW Human;
KW imaging
KW diabeti
KW tumour
KW angioge
KW reverse
XX Homo sa
XX Homo sa
XX O6-NOV-
PR 06-NOV-
PR 07-MAR-
XX IBicknel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a new oligonucleotide label-domain CC comprising the sequence (CTATTTT) n and its complement (AAAATAG) n, where n is 1. The probe sets of the invention are useful for detecting kappa or CC lambda-immunoglobulin light chain mRNA or corresponding heteronuclear RNA, CMV (cytomegalovirus) immediate early RNA, EBV (Epstein-Barr virus) CC early RNA 1 and RNA 2, and human Alu repetitive satellite genomic control on the invention is a useful generic sequence for incorporation into oligonucleotide probes for detecting gene-specific sequences within CC cells or tissue samples in in situ hybridisation analysis and for cells or tissue samples in immunobistochemical analyses. The present mucleic control of the probes that were used in the invention for detecting or oligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or oligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting or colligonucleotide probes that were used in the invention for detecting the colligonucleotide probes to the colligonucleotide probes for the coll
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel oligonucleotide label-domain for incorporation into oligonucleotide probes useful for detecting or localizing nucleic acid target genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK87131 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-371972/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-SEP-2000; 2000US-0233177P
  Bicknell R,
                                                                                                                                                                                                                                                                                                                                                                                                         imaging vascular endothelium; proliferative disease; cancer; psoriasis; diabetic retinopathy; atherosclerosis; menorrhagia; endothelial damage; tumour neovasculature; cardiac disease; endometriosis; hypoxic condition; angiogenesis; cytostatic; RT-PCR; connective tissue growth factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human connective tissue growth factor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-OCT-2002
                                                                                                      06-NOV-2000;
07-MAR-2001;
                                                                                                                                                                               06-NOV-2001; 2001WO-GB004906.
                                                                                                                                                                                                                                                                                   WO200236771-A2
                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; endothelial cell-specific molecule 4; ECSM4; neovasculature;
                                                                                                                                                                                                                                    10-MAY-2002
                                                       (IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tissue sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4463 CTTTTTTTTTTTTTTTTTT 4483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a cell or tissue sample.
                                                                                                                                                                                                                                                                                                                                                                                    transcription-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTATTTTCTATTTTCTTTT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 19; 71pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
  Huminiecki L;
                                                                                                      2000US-0245566P.
2001US-0273662P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Connaughton J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 C; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RT-PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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Claim 1; Page 4;

38pp;

English

Novel common primer useful for diagnosing cancer, is made from highly homologous areas of twelve melanoma antigen gene subtypes and eight G

and eight GAGE

WPI; 2002-026166/03

Park J, (ICGI-)

Jeon

IC & G CO

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 3306
AAD22642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer, psoriasis, diabetic retinopathy, atherosclerosis or menorrhagia. The ECSN4 proteins are also useful in the manufacture of diagnostic or prognostic agent for such conditions. The proteins are also useful for detecting endothelial damage or activation, detecting a tumour or tumour neovasculature, cardiac disease, or endometriosis by detecting the amount of ECSN4 present in a sample. The polynucleotide sequences encoding ECSN4 are useful in gene therapy for treating a hypoxic condition such as cancer, cardiac disease, endometriosis or atherosclerosis and in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to endothelial cell-specific molecule 4 (BCSM4), and the polynucleotide sequences encoding it. The ECSM4 proteins are useful for imaging vascular endothelium in the body of an individual, and for diagnosing and treating a proliferative disease or condition involving the vascular endothelium (preferably, neovasculature) such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel endothelial cell-specific molecule polypeptide 1 or 4, useful imaging, diagnosing and treating a condition involving vascular endothelium e.g. cancer, cardiac disease, endometriosis, diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    manufacture of medicaments for treating the above disease. The sequenare useful for modulating angiogenesis in an individual. The present sequence represents a RT-PCR primer for RNA encoding human connective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 9 A; 4 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD22642 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tissue growth factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 165; 248pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-508120/54.
                                                                                                                                                                                                                                                                                                                                                                                                                           25-APR-2000; 2000KR-00021837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2001; 2001WO-KR000681
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer; melanoma antigen gene; MAGE; GAGE; cancer; reverse transcription polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTTTTCACCTGGAAGCATTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              #3, used as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.6; Db 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnostic marker for MAGE subtypes 1-6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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RESULT 3307
AAD33020/c
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Best Local
                     terminal portion of a coding sequence of an RNA virus and at least an N-terminal portion of a coding sequence of RNA virus, a second polynucleotide region including a 3' untranslated region (UTR) sequence of the RNA virus and at least a C-terminal portion of a coding sequence of the virus and a third polynucleotide region encoding a reporter molecule, flanked by first and second polynucleotide regions; and a promoter sequence being operatively linked to expression cassette in a promoter sequence being operatively linked to expression cassette in a manner so as to enable a transcription of a minus strand RNA molecule from the expression cassette. Nucleic acid construct of the invention is useful for detecting the presence of an RNA virus in a cell. It is also useful for screening anti-viral drugs and determining drug resistance of an RNA virus. The present sequence is a PCR primer used to amplify the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to primers useful for diagnosis of one or more kinds of cancer and a diagnostic kit comprising the primers of the invention. The primers are derived from highly homologous areas of twelve melanoma antigen gene (MAGE) subtypes or eight GAGE subtypes. The primers of the invention are useful for diagnosing cancer by performing polymerase chain reaction (PGR), reverse transcription (RT)-PCR, and nested PCR. The kit of the invention is used to detect six MAGE subtypes and eight GAGE subtypes. The present DNA sequence is an antisense PCR primer which is used as a diagnostic marker for MAGE subtypes 1-6
                                                                                                                                                                                                                                                                                                                                                             The invention relates to nucleic acid expression cassette including a first 5' non-coding region (NCR) sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel nucleic acid construct useful for detecting the presence of RNA virus, comprises an expression cassette and a promoter operably linked expression cassette for minus strand RNA transcription of the cassette.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tan YH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 24; 81pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JUL-2000; 2000US-0220248P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUL-2001; 2001WO-IL000669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   untranslated region; UTR; anti-viral drug; drug resistance; primer; PCR; HCV-S1; Hepatitis C virus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-280605/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200208447-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HCV-S1 overlapping cDNA region amplifying antisense PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD33020 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INST MOLECULAR & CELL BIOLOGY. EHRLICH G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       construct; expression cassette; non-coding region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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the genome
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RESULT 3309 ABK52977

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AAL46645/c
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Best Local S
Matches 17
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                                                    Query Match
Best Local (
                                                                                                                The present invention relates to the use of a nucleic acid that causes the synthesis of the AKINI1 protein, to induce or increase the resistance to pathogen attack in plants. The nucleic acid and its encoded protein can be used to impart resistance to bacteria, viruses, fungi and nematodes, especially necrotropic pathogens such as Xanthomonas campestris, in large-scale crops, vegetables and flowers. Probes and primers that hybridise with the AKINI1 gene can be used to detect resistance against pathogens, and antisense sequences can be used to modulate resistance. The present sequence is a PCR primer used to isolate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                              WPI;
                                                                                  Sequence 21
                                                                                                         the AKIN11
                                                                                                                                                                                                                             Example 2; Page 74; 76pp;
                                                                                                                                                                                                                                                           Inducing or increasing resistance to pathogens in plants e.g. increase flowers and vegetables, by introducing a nucleic acid that
                                                                                                                                                                                                                                                                                                                  Roby D,
                                                                                                                                                                                                                                                                                                                                        (INRG )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fungicide;
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CENT NAT RECH
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GTGGGAGCTCCTCCAAGTTCC
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                                                                                                       coding sequence of the invention
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                                                                                   5 A;
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Pred. No. 2
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Detecting target analyte comprises providing a first classifier of a first population of sensors, distributing a second population of sensors and determining the response of the second population of sensors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; interleukin 6; microsphere; genosensor; target; ss; microarray; optical signature.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK52977 standard; DNA; 21 BP
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genosensor and
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/mod_base= OTHER
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  used to illustrate the
  method of the invention
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Query Match
Best Local Similarity
Matches 17; Conserv

Conservative

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Score 14.6; DB 1; Length 21; Pred. No. 2.3e+03; O; Mismatches 4; Indels

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                                                                                                                          The invention relates to detecting a target analyte (TA) comprising: (a) comproviding first classifier (I) for the response of a first population of sensors (II) to a first population of distributing a second population of sensors (II) to a first TA; (b) condition (c) determining response of (IV) to the sample, where response condition (c) determining response of (IV) to the sample, where response condition (c) determining response of the sample, where response conditions a condition of microspheres comprising an array comprising: (a) providing a contacting the microspheres with a sample of the target analyte; (b) contacting the microspheres with a sample of the target analyte; (c) contacting a classifier for the response of the microspheres on a substrate with carget analyte; and (e) distributing the microspheres on a substrate with a surface comprising discrete sites. The new method detects a target analyte in a sample by contacting the sample with a sensor array. The contactive agents may be synthesised on the beads which are then randomly distributed on a patterned surface. The beads are self-encoded with dyes allowing a correlation of the location of an individual site on the carray. The self-encoding feature eliminates the need for a more complex, the array are self-encoded by exposing the array to a reference analyte the array are self-encoded by exposing the array to a reference analyte.
                                 while illuminating the array with excitation light energy. The light sensor array may carry thousands of discrete sensing elements whose combined signal provides for substantial improvements in sensor dete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting target analyte comprises providing a first classifier of a first population of sensors, distributing a second population of sensors and determining the response of the second population of sensors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 38; 104pp; English.
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The invention relates to detecting a target analyte (TA) comprising: (a) CC providing first classifier (I) for the response of a first population of CC sensors (II) from a first pool of sensors (III) to a first TA; (b) CC distributing a second population of sensors (IV) from (III) on an array; CC resembles (I) for a first TA, indicating the presence of the first TA in CC resembles (I) for a first TA, indicating the presence of the first TA in CC contacting the microspheres comprising an array comprising; (a) providing a CC population of microspheres with a sample of the target analyte; (b) CC recording the response of the microspheres to the target analyte; (d) CC generating a classifier for the response of the microspheres on a substrate with a rarget analyte; and (e) distributing the microspheres on a substrate with a sample by contacting the sample with a sensor array. The method allows the synthesis of the biocactive agents i.e., nucleic acids and rethod allows the synthesis of the biocactive agents i.e., nucleic acids array in the microspheres to the synthesis of the biocactive agents i.e., nucleic acids
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ABK52964/c
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and antibodies, to be separated from their placement on an array. The bioactive agents may be synthesised on the beads which are then randou distributed on a patterned surface. The beads are self-encoded with distributed on a patterned surface. The beads are self-encoded with distributed on a patterned surface. The beads are self-encoded with distributed on a patterned surface.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 19; Fig 22; 104pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; interleukin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.2%;
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Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
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                         randomly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KW Human; ds; cytochrome P450 Al; CYP4501Al; UGT2B4; MDR1;
KW cytochrome P450 A2; CYP4501A2; cytochrome P450 O2E; CYP45002E1; LTF;
KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW dyntathione-5-transferase 12; GST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
KW MADPH quinone oxidoreductase 2; WO2; sulfotransferase thermolabile; STM;
KW WDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7;
KW WDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW multidrug resistance associated protein 3; canner; prostate;
KW multidrug resistance associated protein 3; canner; prostate;
KW multidrug resistance associated protein 3; canner; prostate;
KW acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR4; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW altered largenges associated protein 3 canner; prostates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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                                                                                                                                                          WPI; 2002-698522/75
                                                                                                                                                                                                                                                                                                                       28-NOV-2001; 2001WO-US044838.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 allered urug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human epoxide hydroylase 2 polymorphic sequence #54.
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                                                                                                                                                                                                                                                                               28-NOV-2000; 2000US-00724389
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                                                                                                                                                                                                                                                                                                                                                                                                        WO200257410-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                           (DNAS-)
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                                                                                                                                                                                                                                         DNA SCI LAB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 1 C; 10 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphism.
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Indels
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Isolated nucleic acid molecules having polymorphisms in known human e.g. cytochrome p450 and cathepsin S useful as genetic linkage market for locating, identifying and characterizing the genes responsible if

enetic linkage markers genes responsible for

Example 10; Page 119; 714pp; English

disorder-related

traits.

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RESULT 3313
ABS98273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
Human; ds; Cytochrome P450 Al; CYP4501Al; UGT2B4; MDR1; Cytochrome P450 A2; CYP4501A2; Cytochrome P450 O2E; CYP45002El; LTF; adrenergic receptor betal; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112; aryl hydrocarbon receptor nuclear translocator; ARNI; cathepsin S; CTSS; cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase 12; GST12; histamine-N-methyl transferase; HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT; NNADPH quinone oxidoreductase 2; NOO2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B7; UGT2B7; UDP-glucuronosyl transferase 2B7; multidrug resistance 1; lactotransferain; orphan nuclear receptor; multidrug resistance associated protein 3; cancer; prostate; acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betai (ADBR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                        Human lactoferrin (LTF) gene polymorphic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                            23-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABS98273 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 8.A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA sequence of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
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Pred. No. 2.
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Matches

337 TACTTTGAGGTGGACATCCCT

Loca1

l Similarity 17; Conser

Conservative

<u>,</u>

Mismatches

0.2%; ე წ σ ç,

Score 14.6; Pred. No. 2 Pred.

.3e+03 DB 1;

Length Indels

21;

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Gaps

0

Sequence 21

B₽;

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9

T; 0

U; 0 Other

DNA sequence of

the invention

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CC aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (FLAP), epoxide hydrocarbon receptor nuclear translocator (FLAP), epoxide hydrocarbon receptor nuclear translocator (FLAP), epoxide hydrocarbon receptor nuclear translocator (FLAP), glutathione-S-transferase 2 (ESPHX2), 5-lipoxygenase activating CC protein (FLAP), glutathione-S-transferase 12 (GST12), histamine-N-methyl CC transferase (NNMT), NADH quinone oxidoreductase 2 (NOC2), CC (UGT2B4), UDP-glucuronosyl transferase 2 (NOC2), CC (UGT2B4), UDP-glucuronosyl transferase 2 (NOC2), CC (MRR3), urokinase receptor (URAP), multidrug resistance 2 (NOC2), CC (MRR3), lactotransferrin (LTP), multidrug resistance associated protein 3 (CC (MRR3), orphan nuclear receptor (NR112), or acetylcholine muscarinic CC receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR4 or CHMR5) sequence CC The polymorphisms in the human genes cited in the invention are useful as CC genetic linkage markers for locating and characterising the genes that CC are responsible for specific traits within the genome and eventually CC identifying the genes responsible for a variety of disorder-related CC traits as a result of their e.g., overexpression, constitutive traits and resulting the comprising the comprising the golymorphic sequences contained in CYP4501A; CYPP4501A; CYP74502E1, CC ARM7, EPHX2, GST12, NMT, NOO2, NR112, STM, UGT2B4, UGT2B15, AHR, CM AHR, MDR1 and/or MDR3 are useful for screening individuals for altered drug cmetabolism. The polymorphic sequences contained in CYP4501A; CYPP4501A1, CYPP4501A2, CYPP45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological; SNP; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A1), adrenergic receptor betal (ADBR1), cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guida
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-NOV-2000; 2000US-00724389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-NOV-2001; 2001WO-US044838.
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aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (CDX2), diazepam binding cinhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase activating protein (FLAP), glutathione-S-transferase 12 (GST12), histanine-N-methyl transferase (HNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl transferase (NNMT), NADPH quinone oxidoreductase 2 (NQC2), sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 284 (UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase (UGT2B15), urokinase receptor (URA), multidrug resistance 1 (VMR1), lactotransferrin (LTFP), multidrug resistance associated protein 3 (VMRP3), orphan nuclear receptor (NRAI), or acetylcholine muscarinic creceptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR4 or CHMR5) sequence. The polymorphisms in the human genes cited in the invention are useful as genetic linkage markers for locating and characterising the genes that are responsible for specific traits within the genome and eventually cleantifying the genes responsible for a variety of disorder-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KW Human; 88; primer; cytochrome P450 Al; CYP4501Al; UGT2B4; MDR1; PCR; W Cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002El; LTF; KW cytochrome P450 A2; CYP45002El; LTF; KW adrenergic receptor betal; ADBR1; aryl hydrocarbon; ARR; MRP3; NR112; KW cytlooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; KW cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; W epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; KW HAMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT; KW HAMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT; KW HAMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NTM; W DP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7; W UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; W UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; W UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; W ultidrug resistance associated protein 3; cancer; proteate; wullocation muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5; KW altered drug metabolism; cardiovascular function; colorectal tumour; KW central nervous system; nulmonary; immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP4501ZE1), adrenergic receptor beta1 (ADBR1), cytochrome P450 02E1 (CYP4500ZE1), adrenergic receptor beta1 (ADBR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
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ABS97692 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 13; Page 124; 714pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorder-related traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-698522/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Guida M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   central nervous system; pulmonary; immunological.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Histamine N-methyl transferase (HNMT) polymorphic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-NOV-2000; 2000US-00724389.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-NOV-2001; 2001WO-US044838.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hall J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TACTTTGTGGTGGTCACTCCT 21
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                                                                                                                                                                                                                                                                                                                                                                                        KW cytochrome P450 A2; CytoShrome P450 A1; CYP4501A1; UGT2B4; MDR1;
KW cytochrome P450 A2; CYP4501A2; cytochrome P450 O2E; CYP45002E1; LTF;
KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; CCX2; diazepam binding inhibitor; DBI; haematological;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW glutathione-5-transferase 12, GST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase;
KW MADPH quinone oxidoreductase 2; NQO2; sulforransferase thermolabile; STM;
KW MADPH quinone oxidoreductase 214; UDD-glucuronosyl transferase 2B7;
WUDP-glucuronosyl transferase 2B4; UDD-glucuronosyl transferase 2B7;
KW MUDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW multidrug resistance 1; lactotransferrin; orphan nuclear receptor; uPA;
KW multidrug resistance associated protein 3; cancer; prostate;
KW acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW central nervous system; pulmonary; immunological; sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression, mutation or underexpression, which may be used in diagnosing CC and/or treating the disorders. The nucleic acid molecules comprising the polymorphic sequences contained in CYP4501A1, CYP44502E1, CYP44502E1, STM, UGT2B4, UGT2B4, UGT2B15, AHR, URR1 and/or MDR3 are useful for screening individuals for altered drug metabolism. The polymorphic sequences contained in CYP4501A1, CYPP4501A2, CHR, WDR1 and/or MDR3 may also be used to screen individuals for susceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are used to screen for altered central tumours, in DB1 or CHMR1 for altered central cused to screen function, in CNZ for altered cuseptibility to colorectal tumours, in DB1 or CHMR1 for altered central nervous system function, in FLAP and HNMT for altered pulmonary, communological or haematological function, in KLKZ for altered serine cused control, in CHMR3, CHMR5 for altered central and peripheral nervous system function. The present sequence represents a PCR primer used to amplify the sequences of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human acetyl choline muscarinic receptor 1 polymorphic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 2 A; 6 C; 4 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS98507 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   traits as a result of their e.g., overexpression, constitutive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1295 CCATGATAAAAGGCCACAGCTA 1315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 CCAGGATAAAGAGGÁCAGCTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
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Isolated nucleic acid molecules having polymorphisms in known human

genes

WPI; 2002-698522/75

Guida M,

Hall J;

(DNAS-)

DNA SCI LAB

28-NOV-2000; 2000US-00724389 28-NOV-2001; 2001WO-US044838. 25-JUL-2002.

Homo sapiens.

WO200257410-A2

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CC thuman cytochrome P450 Al (CYP4501Al), adrenergic receptor betal (ADBRI), cc aryl hydrocarbon (AHR), aryl hydrocarbon receptor muclear translocator (CRANT), cathepsin $ (CTSS), cyclooxgenase 2 (COX2), diazepam binding crotein (FTAP), glutathione-5-transferase 12 (CGYI2), histemaine-N-methyl ctransferase (HNMT), kkalikrein 2) kkk2, nicotinamide -N-methyl ctransferase (HNMT), kkalikrein 2) kkk2, nicotinamide -N-methyl croteransferase (HNMT), which pullidry resistance associated protein 3 (CC (MPRI), lateotransferrin (LTP), multidryg resistance associated protein 3 (CC (MPRI), lateotransferrin (LTP), multidryg resistance associated protein 3 (CC (MPRI), lateotransferrin (LTP), multidryg resistance associated protein 3 (CC (MPRI)) orphan nuclear receptor (NRII), or acetylcholine muscarinic creeptor (1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR2, CHMR4 or CHMR5) sequence. CC receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR2, CHMR4 or CHMR5) sequence. CT is polymorphisms in the human genes cited in the invention are useful as genetic linkage markers for locating and characterising the genes that CC are responsible for specific traits within the genome and eventually ctraits as a result of their e.g., overexpression, constitutive expression, mutation or underexpression, which may be used in diagnosing cand/or treating the disorders. The nucleic acid may discompating the genes compating the genes compating the genes compating the genes compating the genes constitutive colorest polymorphic sequences contained in CYP4501A2, CYP4502E1, CC MRT, while may be used to screen individuals for altered drug compating the genes compating 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
Sequence
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     21
     BP;
     Ŋ
A;
7 C; 7 G;
U
T; 0 U; 0 Other;
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RESULT 3316
ABS98319/c
ID ABS9831
XX
AC ABS9831
XC ABS9831
XX
DT 23-DEC-
XX
DE Human 1
XX
Human;
KW Human;
KW drochn;
KW adroch;
KW adroch;
KW aryl hy
KW cycloo
KW cycloo
KW cycloo
KW glutat;
KW HNMT; }
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Best Local Similarity
Matches 17; Conser
Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1; cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF; cytochrome P450 02E; CYP45002E1; LTF; adrenergic receptor beta1; ADBR1; ary1 hydrocarbon; AHR; MRP3; NR112; ary1 hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS; cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase 12; GST12; histamine-N-methyl transferase; HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
                                                                                                                                                                                                           Human lactoferrin (LTF) gene polymorphic sequence #82
                                                                                                                                                                                                                                                                                                                                                   ABS98319 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5161 TTCTCCTGGGACAGTGGGCTC 5181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGCTCATGGGCCACTGGGCTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.2%;
nilarity 81.0%;
Conservative
                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.6; DB 1;
Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
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NADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7; UDP-glucuronosyl transferase 2B7; UDP-glucuronosyl transferase; uPA; multidrug resistance 1; lactotransferrin; orphan nuclear receptor; multidrug resistance associated protesin 3; cancer; prostate; acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5; altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological; SNP; single nucleotide polymorphism.

Homo sapiens.

WO200257410-A2

25-JUL-2002

28-NOV-2001; 2001WO-US044838.

28-NOV-2000; 2000US-00724389

(DNAS-) DNA SCI LAB INC

Guida

Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits.

23; Page 149; 714pp; English.

CC aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (C aryl hydrocarbon (SHR), aryl hydrocarbon receptor nuclear translocator (C ARWT), cathepsin \$ (CTSS), cyclooxgenase 2 (CDX2), diazepam binding conhibitor (DBI), epoxide hydroxylase 2 (EDHX2), 5-lipoxygenase activating cyrotein (FLAP), glutathione-5-transferase 2 (CDX2), histamine-N-methyl curansferase (NNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl transferase (NNMT), MADPH quinone oxidoreductase 2 (NQO2), cransferase (NMT), NADPH quinone oxidoreductase 2 (NQO2), cransferase 2B4 (CC guifotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (CC guifotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (CC guifotransferase (UGT2B15), urokinase receptor (NPA), multidrug resistance associated protein 3 (CMR12), lactotransferrin (LTF), multidrug resistance associated protein 3 (CMR21), lactotransferrin (LTF), multidrug resistance associated protein 3 (CMR21), lactotransferrin (LTF), multidrug resistance associated protein 3 (CMR21), or acetylcholine muscarinic craceptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR4 or CHMR5) sequence. CC monthly are protein for locating enes cited in the invention are useful as a result of the human genes cited in the invention are useful as creenifying the genes responsible for a variety of disorder-related cracity of disorder-related cracity of an acetylcholine muscaring the capture of treating the disorder of traits within the genome and eventually cc and/or treating the disorder of their e.g., overexpression, constitutive useful as a result of their e.g., overexpression, constitutive useful as comprising the genes comprising the genes comprising the genes comprising the genes of their e.g., overexpression, which may be used in diagnosing the genes comprising the genes of their e.g., overexpression, constitutive constitutive and/or mins are useful for screening individuals for altered drug comprising the genes comprising the genes comprising the genes comprising the genes compri This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP45002E1), adrenargic receptor beta1 (ADBR1), polymorphic haematological function, in CHMR3, CHMR4 or CHMR5 for altered central nervous system function. DNA sequence of the invention The present sequence represents a

Sequence 21 BP; в P Ç 10 ဂ္ 2 T; 0 U; 0 Other;

DB 1;

Length

21;

5

i k

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RESULT 3317
ABS97961;
ID ABS97961;
XX Human UDP-glucuronos;
XX Human; ds; Cytochrome P450 A2;;
XX Algrenergiz receptor |
XX adrenergiz receptor |
XX ABS97961;
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KW Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
KW cytochrome P450 A2; CYP4501A2; cytochrome P450 O2B; CYP45002B1; LTP;
KW adrenergic receptor beta1; ADBR1; arryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
KW glutathione-S-transferase; JGST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
KW NADPH quinone oxidoreductase 21, NQO2; sulfotransferase thermolabile; STM;
KW NADPH quinone oxidoreductase 224; UDD-glucuronosyl transferase 2B7;
WUP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW multidrug resistance 1; lactotransferrin; orphan nuclear receptor;
KW multidrug resistance associated protein 3; cancer; prostate;
MCW acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW altered and protein system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 17
This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 O2E1 (CYP45002E1), adrenergic receptor betal (ADBR1), aryl hydrocarbon (AHR), aryl hydrocarbon receptor muclear translocator (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding inhibitor (DBI), epoxide hydroxylaxee 2 (EPHX2), 5-lipoxygenase activating protein (FLAP), glutathione-S-transferase 12 (GST12), histemine-N-methyl transferase (NNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl transferase (NNMT), NADPH quinone oxidoreductase 2 (RQC2), sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl transferase 2B4 transferase (UGT2B15), urokinase receptor (uPA), multidrug resistance 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated nucleic acid molecules having polymorphisms in known human genes. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 20; Page 137; 714pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-NOV-2000; 2000US-00724389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorder-related traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-698522/75.
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Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the disease.

Brinkmann

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Hoffmeyer

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19-SEP-2000; 2000EP-00120123 31-AUG-2001; 2001WO-EP010087 WO200224747-A2

28-MAR-2002.

single nucleotide polymorphism; PCR; primer; ss

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RESULT 3318
ABK94085/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                           Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK94085 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Endothelin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EDN-1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP detection PCR primer
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Claim 1; Page 55; 190pp; English.

describes a polynucleotide (I) of the endothelin

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Novel polynucleotide of the endothelin/endothelin converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP;
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                                                                                       WPI; 2002-435060/46.
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                                                                                                                                                                                                                                                                  (EPID-) EPIDAUROS BIOTECHNOLOGIE
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17; Conserv
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                                                                                                                                                                      Hoffmeyer
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CC signaling system which is associated with a cardiovascular disease (I), CC the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) CC or (II) is useful for producing cells capable of expressing a molecular CC variant polypeptide which is associated with a cardiovascular disease. CC (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing and complecular variant gene comprising (I) is useful for identifying and cobtaining a pro-drug or drug capable of modulating the activity of a cC molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system (CC or its gene product, or for identifying and obtaining an inhibitor of the cC signaling system or its gene product. The isolated proteins and CC polynucleotides encoding them are useful for preparation of a phypercholesterolacial composition for treating a cardiovascular disease such as CC coronary heart disease, hypertension, atherosclerosis, or related to hypercholesterolacmia. The gene or a polynucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or dentify single nucleotide polymorphisms in DNA encoding primer used to identify single nucleotide polymorphisms in DNA encoding pathway XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a polynucleotide (I) of the endothelin (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (signaling system which is associated with a cardiovascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful for treating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 55; 190pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
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Query Match Best Local Matches

0.2%;

Score 14.6; Pred. No. 2. ed. No. 2.3 Mismatches

2.3e+03;

4.

0

Gaps

0

21 BP; 3

A; 3 C; 3

G; 12 T; 0 U; 0 Other;

DB 1; Length 21;

0

Sequence 21 BP; 12 A; 3 C; 3 G;

3 T; 0 U;

0 Other;

CC (EDN) endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) CC signaling system which is associated with a cardiovascular disease. (I) CC the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) CC or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (C (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing a complex variant gene comprising (I) is useful for identifying and complex variant of a polypeptide of modulating the activity of a cell expressing a considerable reproduct, or for identifying and obtaining a pro-drug or drug capable of modulating the activity of a cell-expressing a considerable reproduct, or for identifying and obtaining an inhibitor of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and comparison of the encoding them are useful for preparation of a polypeptide of the EDN/EDNR/ECE coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial C hypercholesterolaemia. The gene or a polymucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polymucleotides, genes, vectors, polypeptides, antibodies or dentify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway

문 ş Query Match Local 3722 TCCTCATTCATTGAGCTTTTT 3742 _ l Similarity 17; Conserv TCCTGATTTAGTGATCTTTTT 21 Conservative 81.0%; 0.2%; 0, Score 14.6; Pred. No. 2. Mismatches .3e+03 DB 1; Length 21; Indels 0, Gaps 0

RESULT 3320 ABK94081/c ID ABK9408 XAXRXRXRXRXRXRXRXRXRXRXRXRXRXX 27-AUG-2002 ABK94081; ABK94081 standard; DNA; 21 (first entry) ВP

Endothelin-1 (EDN-1) SNP detection PCR primer #25.

Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; single nucleotide polymorphism; PCR; primer; ss. Synthetic 31-AUG-2001; 2001WO-EP010087 28-MAR-2002 WO200224747-A2 2000EP-00120123

Brinkmann U,

Hoffmeyer S

(EPID-) EPIDAUROS BIOTECHNOLOGIE

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ABK94242
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Best Local (
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                                                                                                                                                                                                                                                                                                        hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP;
                                                                                                                                                                                                                                                                                                                                                                                  Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprising polynucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
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                                                                                           31-AUG-2001;
                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                       single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK94242 standard; DNA;
                                             19-SEP-2000; 2000EP-00120123
                                                                                                                                              28-MAR-2002
                                                                                                                                                                                         WO200224747-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         converting enzyme 1 (ECE-1) SNP detection
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                                                                                           2001WO-EP010087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                       polymorphism; PCR; primer;
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Pred. No. 2.
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The invention describes a polynucleotide (I) of the endothelin CC (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) CC signaling system which is associated with a cardiovascular disease. (I), CC the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) cor (II) is useful for producing cells capable of expressing a molecular CC variant polypeptide which is associated with a cardiovascular disease. (C (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing and CC molecular variant gene comprising (I) is useful for identifying and CC molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system CC or its gene product, or for identifying and obtaining an inhibitor of the CC activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system CC signaling system or its gene product. The isolated proteins and CC polynucleotides encoding them are useful for preparation of a CC coronary heart disease, hypertension, atherosclerosis, or related to CC abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polynucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for CC comprising polynucleotides, genes, vectors, polypeptides, antibodies or bost cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding the polymorphisms in DNA encoding
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    Sequence 21
                                                                                              cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
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A; 4 C; 4 G; 8 T;
0 U;
         0 Other;
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Matches
ABK94082 standard; DNA;
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                                                                  l Similarity
17; Conser
                                                  GTTTTCAGCTGGGGCTTGAAA 5375
                                                                  Conservative
                                                                        0.2%;
21
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Local

Score 14.6;

DB 1; Length 21;

0

Endothelin-1 (EDN-1) 27-AUG-2002 ABK94082 (first entry) SNP detection PCR primer #26.

hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP; Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; Synthetic nucleotide polymorphism; PCR; primer;

31-AUG-2001; 2001WO-EP010087

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ABN86025
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 4 A; 3 C; 2 G; 12 T; 0 U; 0 Other;
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   02-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mutagenic primer D399S
                                                                                                                                                    23-MAY-2002
                                                                                                                                                                                                                                                                                                    Synthetic
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                                                                            23-MAY-2001; 2001US-00863693.
                                                                                                                                                                                                                            US2002062010-A1
                                                                                                                                                                                                                                                                                                                                                                             antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                           Antibody; bispecific antibody; immunoadhesin; cytostatic; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABN86025;
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                     vaccine; tumour;
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       97US-0046816P
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                                                                                                                                                                                                                                                                                                                                                                 PCR primer; 88.
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Pred. No. 2.
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The invention relates to a new multispecific antibody, comprising at CC least two polypeptides (PP1 and PP2) which meet at a multiface, where PP1 chas a multimerisation domain forming an interface positioned to interact with an interface of a multimerisation domain of PP2, and both CC with an interface of a multimerisation domain of PP2, and both cCC avariable light chain, where the light chain has a sequence common to CC artibodies. Hetromultimers of the inventions include bispecific CC antibodies, bispecific immunoadhesins and antibody-immunoadhesin cC chimeras. The activity of antibodies of the invention may be described CC as, cytostatic, antibacterial and antivital. The heteromultimer can be cused for redirected cytotoxicity, for example to kill tumour cells, as a converting enzyme activated prodrugs at a target site such as a tumour, CC converting enzyme activated prodrugs at a target such as a tumour, CC for treating infectious diseases, for targeting immune complexes to cell current sequence represents the mutagenic primer D399S for introducing a cutation into the CH3 domain of a humanised anti-CD3 heavy chain or CD4-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New multispecific antibodies having heteromultimeric and common components are useful to direct treatment to a target site such as a tumor cell, cell surface receptor or clot, as a vaccine adjuvant and treat infectious disease.
Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 23; 36pp; English
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                                          site directed mutagenesis
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BP; 3
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7 C; 9
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  G; 2 T; 0 U; 0 Other;
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밁 Ş Query Match
Best Local Similarity
Matches 17; Conserv 185 GCCGCCTGACCTCCGCACGGG 205 GCCGTCGGAGCTCAGCACGGG Conservative 0.2%; 0, 21 Mismatches DB 1; Length 4 Indels 21; 0 Gaps

0

RESULT 3324 AAL49183/c 30-OCT-2002 AAL49183; AAL49183 standard; DNA; 21 BP (first entry)

CD 151; porcine reproductive and respiratory selective breeding; xenotransplant; anti-RNA anti-viral; vaccine; PCR; primer; 88. syndrome virus; PRRSV; pig; entry protein; anti-REP;

Porcine CD 151 coding sequence PCR primer #7.

Sus scrofa.

WO200260924-A2

08-AUG-2002.

29-JAN-2002; 2002WO-US002868

29-JAN-2001; 2001US-00772044. 28-JAN-2002; 2002US-00772044.

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Kapil S, Shanmukhappa K;

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ABZ95973/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC The present invention relates to a method of determining the susceptibility or resistance of an animal to porcine reproductive and crespiratory syndrome virus (PRRSV). This involves assaying for CD 151 in CC addition, coding sequences of CD 151 ere described, and anti-viral CC compounds designated anti-RNA entry proteins (anti-REPs). The method is CC useful for determining susceptibility and resistance to PRRSV in an CC animal. This is particularly useful for improving swine breeding or for CC screening different pig breeding lines. The method is also useful for CC caping non-simian recombinant cell lines for propagating the virus, CC against pRRSV, and for diagnosing pRRSV infection in a swine. The present CC sequence is a PCR primer used to isolate the porcine CD 151 coding sequence. Note: The sequence data for this patent did not form part of CC the printed specification, but was obtained in electronic format directly CC from WIPO at fip.wipo.int/pub/published_pct_sequences
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Best Local S
Matches 17
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                                                           Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                              Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining susceptibility and resistance to porcine reproductive and respiratory syndrome virus (PRRSV), useful for improving swine breeding, by assaying for CD 151 in a sample of cellular material of known origin from the animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ95973 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                             WPI; 2003-229219/22
                                                                                                                                                                                                        31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                             Human fibronectin antisense fragment no.1833.
                                                                                                                                                                                                                                                                                                                                                                                                                                                17-OCT-2003
                                                                                                                                          24-APR-2001; 2001US-0286137P
                                                                                                                                                                       23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                    WO200285308-A2
                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 17; Page 35; 77pp + Sequence Listing; English
                                                                                                          (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7406 GCAACATCAGCAGCAGCAGCA 7426
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                                                                                                          EPIGENESIS PHARM INC.
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Tang
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                                                            Sandrasagra A,
L, Shahabuddin
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Pred. No. 2.3e
0; Mismatches
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                                                                           'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  U; 0 Other;
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                                                                           Pabalan
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The invention relates to a novel pharmaceutical composition, which has a Cfirst active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory, antiallergic, antiasthmatic, hypotensive, acc use in antisense gene therapy. The composition is useful for treating or CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels of of or reducing sensitivity to adenosine, reducing levels of adenosine CC using inflammation, lung allergies, or a respiratory disease or condition, CC lung inflammation, but was obtained in electronic format directly from WIPO CC at fig. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       corresponding ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 11215; 872pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               respiration, has oligo(s) antisense to specific gene(s) or it
curresponding RNAs, and glucocorticoid or non-glucocorticoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       steroid
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Query Match Best Local S Matches 17 Sequence 3384 CCTCCCCCAGCTGCCACCCCC 3404 21 l Similarity 17; Conserv CCGCCCACAACGGCCACCCCC 1 21 Conservative BP; 0 A; 0.2%; 3 C; 14 G; 4 T; 0 U; 0 Other; 0, Score 14.6; Pred. No. 2 Mismatches 2.3e+03; DB 1; Length Indels 0 Gaps

0

RESULT 3326 ABX94380 ABX94380; ABX94380 standard; DNA; 21

18-JUN-2003

(first entry)

Human endothelial cell differentiation gene-4, Edg-4 PCR primer #1.

Human; ss; PCR; primer; Edg-4; prostatic disease; Edg-7 receptor; benign prostatic hyperplasia; intracellular signal transduction; endothelial cell differentiation gene; prostate cancer; LPA; lysophosphatidic acid.

Homo sapiens

WO2003013605-A1

20-FEB-2003.

07-AUG-2001; 31-JUL-2002; 06-AUG-2002; 2002WO-JP008016 2001JP-00239306 2002JP-00224215

(NISB) JAPAN TOBACCO INC

Furuno M, Naito T, Yamamoto Aoki ŗ Arai

Kakehi

2003-248240/24

prostatic Composition interaction hyperplasia and on between lysoph for preventing venting or treating prostatic diseases ia and prostate cancer containing inhibiting inhibiting the recommendation of the recommendat its receptor inhibitors of ø ដ

Pharmaceutical composition for treating ailments associated with impaired

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ABX90522/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or inhibiting progression of prostatic diseases, e.g. benign prostatic hyperplasia or their accompanying diseases, comprises substances with activity of inhibiting intracellular signal transduction induced by a stimulus mediated by Edg-7 (endothelial cell differentiation gene-7) receptor (interacting with LPA, lysophosphatidic acid), and pharmaceutically-acceptable carriers. The remedies are for prostatic diseases e.g. benign prostatic carriers are remedies are for prostatic diseases e.g. benign prostatic hyperplasia and prostate cancer. The present sequence is a PCR primer for human Edg-4 used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                 31-JAN-1997;
30-JAN-1998;
19-JAN-2000;
19-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prevent
                         The invention relates to a composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor (VEGF). The antisense oligonucleotide is useful for preparing a composition treating cancer, neoplastic proliferation, abnormal cellular proliferation and preventing angiogenesis. The present sequence is a VEGF CDNA target region for the antisisense oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       angiogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense; ss; human;
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                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-2001;
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                                                                                                                                                                      New composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor, useful for preparing a composition
                                                                                                                                                                                                                                                                         (GILL/)
(MASO/)
                                                                                                                             Example 13; Fig 17B; 54pp; English.
                                                                                                                                                                                                                  WPI; 2003-255224/25
                                                                                                                                                       for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention relates to drug compositions for preventing, treating
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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2000US-00487023.
2001WO-US000019.
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         The present invention describes DNA (I) encoding a corticotropin CC releasing hormone receptor type 1 (CRH-R1) protein comprising an amino CC (II) capable of expressing (I) or its degenerate variant, and regulatory elements necessary for CC (II) capable of expressing (I) or its degenerate variant, and regulatory elements necessary for CC expression of the DNA in a cell; (2) a host cell (III) transfected with CC (II); (3) an isolated CRH-R1 protein (IV) encoded by (I); (4) an antibody CC (v) directed against (IV); (5) a pharmaceutical composition (VI) cC comprising (IV), and a carrier; and (6) protecting (M) skin cells against CC damage induced by an environmental factor, by inducing the expression of CC CRH-R type Ig in the skin cells, where the expression of the receptor protects the skin cells against the damage. CRH-R1 has antipsoriatic, CC antiallergic, immunosuppressive, antiinflammatory and dermatological cativities. (VI) can be used for treating a pathophysiological state such CC as hyperproliferative epidermal disorder, neuroendocrine disorder, altergic contact dermatitis, autoimmune disorder, epidermal constant dermatitis, autoimmune disorder, epidermal constant cardingenesis, malignant transformation of epidermal or dermal constant on chromosome 17, and mouse CRH-R1 is located to chromosome CC is used in an example from the present a PCR primer for mouse CRH-R1, which
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse; corticotropin releasing hormone receptor type 1; CRH-R1; antiporiatic; antiallergic; immunosuppressive; antinflammatory; dermatological; pathophysiological state; neuroendocrine disorder; hyperproliferative epidermal disorder; allergic contact dermatitis; autoimmune disorder; epidermal carcinogenesis; malignant transformation; epidermal melanocyte; chromosome 11; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel DNA encoding corticotropin releasing hormone receptor type 1 which is useful for treating pathophysiological state such as inflammatory skin disease e.g. psoriasis and allergic contact dermatitis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-SEP-2001; 2001US-0322195P
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                                                                                                                                                                                                                                                                                                                                                                                                                 Page 26; 110pp; English.
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Pred. No. 2
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                                                                                                                                                                   The invention comprises a method for inferring a genetic pigmentation trait of a human. The method involves identifying a single nucleotide polymorphism (SNP) in a pigmentation gene - where the pigmentation gene is not melanocortin-1 receptor (MCIR) and agout signaling protein (ASIP). The method of the invention is useful for inferring a genetic pigmentation trait of a human, especially for inferring the race of a human subject. The method is useful for inferring a genetic pigmentation trait such as hair shade or colour, or eye shade or colour of a human subject. The method may be used as a forensic tool for obtaining information relating to physical characteristics of a potential crime victim or a perpetrator of a crime from a nucleic acid sample present at a crime scene. The present PCR primer is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-2001;
26-OCT-2001;
15-NOV-2001;
02-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inferring genetic pigmentation trait such as hair/eye color or shade nucleic acid sample of human subject, by identifying a pigmentation-related haplotype allele of a pigmentation gene in the sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JUN-2001;
07-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; single nucleotide polymorphism; SNP; ss; melanocortin-1 receptor; genetic pigmentation trait; MCIR; agouti signaling protein; ASIP; race; hair colour; eye colour; forensic tool; PCR; primer.
                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-239091/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-MAY-2002; 2002WO-US016789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABT34056 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Frudakis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DNAP-) DNAPRINT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABT34056;
Local Similarity
nes 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pigmentation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GTGCGCCCTGCCATGTGGAAG 843
                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 246; 396pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; 2001US-0300187P.
; 2001US-0310781P.
; 2001US-0323662P.
; 2001US-0344418P.
; 2001US-034674P.
; 2002US-0346303P.
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                                                                                                    B₽;
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001US-0293560P.
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                                                                                                    7
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                                                                                                    A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 6 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
                       81.0%;
                                                                                                    3 C; 8
                                                  0.2%;
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0;
                                                                                                    G; 3 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 3 T; 0
                          Score 14.6; DB 1;
Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.6;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U; 0 Other;
                                                                                                    U; 0 Other;
                       2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID
                                            Length 21;
Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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Gaps

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RESULT 3331
ACF64056/c
ID ACF6405
XX
AC ACF6405
XX

ACF64056 standard;

DNA;

21

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ACF64056

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RESULT 3330
ABX10723/c
ID ABX10723/c
XX ABX1072
XX ABX1072
XX ABX1072
XX IS-APR-
XX Human 9
XX Human 9
XX Homo sa
XX US20021
XX IS-APR-
PF 30-AUG-
XX 20-APR-
PR 20-APR-
XX (HOLL/)
PA (WEBS/)
PA (WEBS/)
PA (THAY/)
XX WPI; 20
XX WPI;
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                                                                                                          Matches
                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                            The invention relates to an isolated glycoprotein hormone Zlutl sequence, the mature protein or antigenic peptides derived from Zlutl. Also included are an isolated polynucleotide encoding Zlutl, an isolated antibody that specifically binds to Zlutl, treating hyperthyroidism in female mammals by administering Zlutl and a pharmaceutical composition comprising Zlutl. Zlutl is useful for manufacturing a medicament for treating hyperthyroidism. Anti-Zlutl antibodies can be used to detect Zlutl in tissue sections from a biopsy specimen or to screen biological samples in vitro for the presence of Zlutl. Zlutl is useful for treating women with hyperthyroidism. The nuclei cacid molecules are useful for detecting the expression of a Zlutl gene in a biological sample. The present sequence is a PCR primer used to detect expression of Zlutl in pituitary cells, amplifying part of the 3' untranslated region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        村
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                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New Zlutl polypeptides and polynucleotides, useful for manufacturing medicament for treating hyperthyroidism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-209228/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Holloway JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-APR-2000;
20-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-AUG-2001; 2001US-00943388
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABX10723 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HOLL/) HOLLOWAY J L. (WEBS/) WEBSTER P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-OCT-2002.
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                                                   4476
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                                                                                                          17;
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                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8; Page 48;
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                                                TTTTTTTGTCTTGAGACATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GATGAAGAGCGTGCTGAAGAC
TTTTCTTTGCCTTGAGTCTTG
                                                                                                                                                                                                                 B₽;
                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0199498P.
2001US-00839706.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zlut1; glycoprotein hormone; hyperthyroidism; romosome 14q23.3; primer.
                                                                                                                                                                                                                 12 A; 4 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hormone
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                                                                                                                              0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thayer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zlutl
                                                                                                       ٥,
                                                                                                                                 Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21
                                                                                                          Mismatches
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                                                                                                                                 DB 1;
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                                                                                                       Gaps
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RESULT 3332
ACF64052/c
ID ACF6405
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ACF6405
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AC ACF6405
ACF
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated polynucleotide (PN) comprising: (a) a sequence comprising at least 15 contiguous nucleotides of a sequence comprising variant sequences (A) from Table 4 given in the specification; or (b) a sequence that is complementary to (A). Also described: (1) an array of (PN) s comprising two or more of the isolated (PN)s; (2) detecting a (PN) in an individual; (3) a computer-readable storage medium, where each record has a field identifying a base occupying a (PN) site and a location of the polymorphic site; and (4) a signal carrying data for access by an application program having executed on a data processing system. The (PN) can be used for detecting loci associated with multiple sclerosis. ACF64025 to ACF64024 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; detection; computer-readable storage medium; polymorphic site; signal carrying data; data processing system; multiple sclerosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-AUG-2001; 2001US-0310741P.
24-SEP-2001; 2001US-0324790P.
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Synthetic.
                                                                                                                                 Human; detection; computer-readable storage medium; polymorphic site; signal carrying data; data processing system; multiple sclerosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 10 A; 7 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polynucleotide, useful for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-268196/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2002; 2002WO-US025268
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     WO2003014319-A2
                                                                                                                                                                                                                                                                                                       13-OCT-2003
                                                                                                                                                                                                                                                                                                                                                             ACF64052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequences used in the exemplification of the present invention
                                                           Synthetic
                                                                                                                                                                                                                                                  ESR1 reverse PCR primer #28.
                                                                                                                                                                                                                                                                                                                                                                                                               ACF64052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DNAS-) DNA SCI INC.
                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6449 CAGTGTTTTTGGATACTTTTT 6469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGGTGTTGTTGGATGCTTTAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 10; 93pp; English
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                                                                                                                                                                                                                                                                                                       (first
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                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                       entry)
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Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                loci associated with multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ΠĽ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Natsoulis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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RESULT 3333
AAL53918
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                                                                                                                                                                                              Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                        signal carrying data for access by an application program having executed on a data processing system. The (PN) can be used for detecting loci associated with multiple sclerosis. ACF64025 to ACF64424 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polynucleotide, useful sclerosis.
                                                                                                                                                                                                                                                             Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jones
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24-SEP-2001; 2001US-0324790P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2002;
18-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 10; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-268196/26.
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                              AAL53918;
                                                         AAL53918 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DNAS-) DNA SCI INC
                                                                                                                                                           5328 CTCTCTTTGCCTCACTCTCTC 5348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HB,
                                                                                                                                     21 CTCTCCTCACCTCACTCCCTC 1
                                                                                                                                                                                              1 Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xu H,
                                                                                                                                                                                                                                                           BP; 6 A; 0 C; 13 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002WO-US025268
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    White R,
                                                                                                                                                                                                              0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for
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                                                                                                                                                                                                                Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detecting
                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            loci associated with multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jin W,
                                                                                                                                                                                                                             Length
                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Natsoulis
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25-MAR-2002; 2002WO-US009126
                                                     03-OCT-2002.
                                                                       WO200277291-A1
                                                                                                         modified_base
                                                                                                                                      Unidentified
                                                                                                                                                       detecting environmental pollutant; pregnancy test; target; ss.
                                                                                                                                                                 enzymatic
                                                                                                                                                                         Target analyte;
                                                                                                                                                                                          Interleukin 6 target sequence.
                                                                                                                                                                 reaction;
                                                                                                                                                               array; substrate; microsphere; non-cleavable linker;
ion; forensic DNA fingerprinting; pesticide; herbicid
                                                                                                            Location/Qualifiers
                                                                                         /*tag= a
/note= "Nucleotide is modified by fluorescein"
                                                                                                                                                                 herbicide;
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(TUFT) TUFTS COLLEGE 23-MAR-2001; 2001US-00816651

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RESULT 3334
ACC42623/c
ID ACC42623 standard; DNA;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel method for detecting a target analyte in the sample. The novel method comprises providing an array comprising an array substrate, at least first and second sites, and a population of microspheres, where first and second reaction components are attached with a non-cleavable linker to the first and second microspheres that are randomly distributed on the sites. The methods are useful for detecting the presence of a particular target analyte, or for detecting an enzymatic reaction, such as the presence or absence of, or mutations on a particular nucleotide sequence or protein, e.g. an enzyme, antibody or antigen. They are also useful in forensic DNA fingerprinting, detecting environmental pollutants such as pesticides or herbicides, and in bacterial, fungal, protozoal, Mycoplasmal, Rickettsial diagnostic tests, as well as in pregnancy tests. This polymucleotide represents a target sequence of the microsphere genosensors of the invention
                                                                                                                                                                                                                                                                                                                                             Human; PCR; primer; transgenic mouse; lymphocyte maturation; IL-3; IL-7; cytokine; interleukin-3; interleukin-6; IL-6; interleukin-7; M-CSF; SCF; macrophage-colony stimulating factor; stem cell factor; oncostatin M; OM granulocyte-colony stimulating factor; GM-CSF; LIF; leukaemia inhibitory factor; ss.
New recipient mammal, preferably a mouse, useful as a model of human disease to assess efficacy of therapeutic or prophylactic treatments, for facilitating production of donor-specific functional immunity.
                                                                                                                                                                                                            05-AUG-2002; 2002WO-US024807.
                                                                                                                                                                                                                                               06-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 28; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         providing an array comprising and microspheres.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting a target analyte or an enzymatic reaction useful for forensic DNA fingerprinting or detecting environmental pollutants, comprises providing an array comprising an array substrate, first and second sites,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Walt
                                                                                                                                                                            23-AUG-2001; 2001US-00938689
                                                                                                                                                                                                                                                                               WO2003018744-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACC42623;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin-6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17;
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                                                                                                                                          GENENCOR INT INC.
                                                                                                      FA, Huang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AATAACCACCCCTGACCCAAC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGAAATACCCCAGACCCAAC 3673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 BP; 8 A; 10 C; 1 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  IL-6, PCR primer hIL6-F2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4:
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ABT33411
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XX
AC ABT3
XX
AC ABT3
XX
V11
DE NOV
XX
W Heps
XW Heps
XW Ant:
KW Ant:
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Best Local S
Matches 17
19-JAN-2001;
23-JAN-2001;
24-JAN-2001;
25-JAN-2001;
25-JAN-2001;
25-JAN-2001;
30-JAN-2001;
02-MAR-2001;
14-MAR-2001;
14-MAR-2001;
15-MAR-2001;
20-MAR-2001;
20-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           macrophage-colony stimulating factor (M-CSF), granulocyte-colony, stimulating factor (GM-CSF), stem cell factor (SCF), leukaemia inhibitory factor (IIF) and oncostatin M (OM). The gene disruption is in a gene that modulated VDJ recombination e.g. a RAG gene. The gene is disrupted by insertion of a transgene comprising major histocompatibility complex (MHC) Class II DR3 and DQ2 genes. The transgenic mouse is useful as a model of human disease to assess efficacy of therapeutic or prophylactic treatments, or to assess the antigenic potential of compounds. The transgenic mouse is also useful for supporting donor haematopoietic stem cells or facilitating production of donor-specific functional immunity. PCR primers ACC42571-ACC42639 were used to generate the transgenic mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 36; 70pp; English.
The present invention relates to a new transgenic mouse, which comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatotropic; immunosuppressive; cardiant; hypertensive; tranquilizer; vulnerary; virucide; antibacterial; protozoacide; fungicide; nootropic; antiparasitic; neuroprotective; cerebroprotective; antiparakinsonian; anticonvulsant; antiaddictive; analgesic; dermatological; keratolytic; antiseborrheic; antirheumatic; antiarthritic; antiinflammatory; anti-HIV; cytostatic; antiasthmatic; antipsoriatic; hypotensive; osteopathic; cytostatic; antiathmatic; antiinfertiality; hypotensive; osteopathic; neuroleptic; antidepressant; antiinfertiality; NOVX; human disease; novX-associated disorder; trauma; viral; bacterial; fungal; protozoal; parasitic infection; Alzheimer's disease; stroke; forensic biology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200281517-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABT33411;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a disruption in both alleles of a gene such that lymphocyte maturation does not occur and exogenous cytokines. The cytokines are selected from interleukin-3 (IL-3), interleukin-6 (IL-6), interleukin-7 (IL-7),
                                                                                                                                                                                                                                                                                                                                                                                                                   22-JAN-2002; 2002WO-US002064.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NOVX PCR primer SEQ ID No 266.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunogen; non-human transgenic animal;
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il Similarity 81.0%;
17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCAACACCAGGAGCAGCCCCA 1
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                              2001US-0264478P.

2001US-0263351P.

2001US-0272870P.

2001US-0275927P.

2001US-0275449P.

2001US-0276449P.

2001US-0277358P.
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                                                                                                                                                                                                                                    2001US-0263799P.
2001US-0264117P.
2001US-0264139P.
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                                                                                                                                                                                                                                                                                                                             2001US-0263598P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy; PCR; primer;
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08-JUN-2001;
10-JUL-2001;
10-JUL-2001;
12-JUL-2001;
09-AUG-2001;
13-AUG-2001;
16-AUG-2001;
18-CCT-2001;
29-NOV-2001;
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20-APR-2001;
20-APR-2001;
30-APR-2001;
17-MAY-2001;
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Vernet CAM,
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Anderson D, Ballinger R, Gerlach V, Spy
, Guo X, Zerhusen B, Andrew D, Mezes
CE, Eisen A, Wolenc A, Baumgartner J, Sh
AM, Taupier RJ, Pena C, Shenoy S, Li L,
B E, Smithson G, Malyankar U, Taillon B,
                                                                                                                                                                                                                                                                                                                                                                                                          2001US-0279857P.
2001US-0285141P.
2001US-0285141P.
2001US-0285141P.
2001US-0287484P.
2001US-0296960P.
2001US-0304353P.
2001US-0304355P.
2001US-0304865P.
2001US-0311289P.
2001US-0311289P.
2001US-03112937P.
2001US-0334198P.
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Mezes P, Patturajan
tner J, Shimkets RA,
S, Li L, Casman S,
                                                                                                                                                                                                                                                                                                                                                                                 Tchernev V,
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WPI; 2003-058504/05. New polypeptides, infections, neurol immunological, skin, renal, brain, muscle and autoimmune disorders tides, designated as NOVX, useful for diagnosing and treating neurological diseases, cancer, allergy, and bone, Liu X; Gusev V; Boldog F;

Zhong

, H; Rastelli

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Example 1; Page 215; 672pp; English.

CC variant differ from the mature form, provided that the variant differs in come more than 15 % of the amino acids from the sequence of the mature cc form. The NOVX polypeptides, nucleic acids encoding the polypeptides, and cc NOVX-associated disorder in humans and for treating a syndrome associated cc with a human disease (NOVX-associated disorder). NOVX polypeptides and cc NOVX-associated disorder in humans and for treating a syndrome associated cc with a human disease (NOVX-associated disorder). NOVX polypeptides and cc predisposition to a disease associated with altered level of polypeptide ce expression or the amount of nucleic acid from a mammal and comparing it collypeptide and polymucleotide, by measuring the level of polypeptide ce expression or the amount of nucleic acid from a mammal and comparing it collypeptide is also useful for identifying an agent that binds to NOVX cc and a cell expression or activity of NOVX. The antibodies and a cell expression or activity of NOVX. The antibodies and a cell expression with presence or amammal for disease. NOVX cc polypeptides are useful for treating a pathological state in a mammal. The antibodies are also constituted in the presence or amount of NOVX in a sample. NOVX cc polypeptides are useful for treating or preventing disorders or syndromes considered to the collypeptide and also treat disorders such as e.g., Alzheimer's cl including trauma, viral, bacterial, fungal, protozoal, and parasitic cc and movx gene and to modulate NOVX encoding nucleic acids are useful for dentifying an agent the NOVX proteins, to detect NOVX mRNA, or a genetic lesion in cc antibodies which are useful for the sequences from very small biological samples such as tissues e.g. that or skin or body fluids in forensic biology and as primers and probes cfor use in identifying and/or cloning NoVX mucleic acids are useful for useful for producing non-human transgenic animals which are useful for cuseful for evaluating modulators of NOVX protein activity. This polymcleotide sequence The invention relates to a novel isolated polypeptide, designated (NOV1 - 33), consisting of a mature form of one of 61 sequences, the specification, or its variant, where amino acid residue(s) in represents a XVON PCR gene therapy. T R primer of the invention designated ä given ä

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Sequence
BP; 4
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U; 0 Other
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The invention describes an isolated POLYX (POLY1-17) polypeptide and its variant. POLYX polypeptides (especially POLY5, POLY5 and POLY7), the polypurcleotides encoding them (I) and an anti-POLYX-antibody (III) are useful for treating or preventing a pathology associated with POLYX

Example

Page

74;

91pp; English.

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RESULT 3336
ABX56493/c
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Best Local
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20-APR-2000;
25-APR-2000;
26-APR-2000;
26-APR-2000;
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09-JUN-2000;
03-JUL-2000;
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11-AUG-2000;
27-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tourette's syndrome; amyotrophic lateral sclerosis; head trauma; Alzheimer's disease; alcoholism; vigilance; anxiety; muscle tension; epileptogenic activity; memory; caratiomyopathy; cancer; angiogenesis; arrhythmogenic right ventricular dysplasia; renal disease; diabetes; Epidermal growth factor like protein; leukaemia; lupus; anaemia; ulcer; haematopoietic stem and progenitor cell like protein; cirrhosis; sulfotransferase-like protein; cholangitis; hepatitis; hyperthyrodism; developmental disorder; Syntaxin-like protein; myxoid liposarcoma; asthma; Lambert-Eaton myasthenic syndrome; acute myeloidleukaemia;
                                                                                          Novel polypeptides and nucleic acids which are members of epidermal growth factor, complement receptor families for diagnosing and trea psychiatric conditions, depression, stroke, Alzheimer's and Parkins
                                                                                  psychiatric disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gamma-aminobutyic acid receptor-like protein; depression; stroke;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABX56493 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transgenic animal; PCR; primer; ss.
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(MACD/)
(SMIT/)
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) ELLERMAN K.
) MACDOUGALL (
) SMITHSON G.
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2000US-0198475P.

2000US-0199476P.

2000US-019980P.

2000US-0200024P.

2000US-02200025P.

2000US-0210809P.

2000US-0215855P.

2000US-0224610P.

2000US-0224610P.

2000US-0271814P.
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Pred. No. 2
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                                                                                                        for diagnosing and treating
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                                                                                               and Parkinson's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease;
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Matches
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20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                       antiinflammatory; plant; antiviral; vasotropic; neuroprotective; cytostatic; cardiovascular; immunosuppressive; respiratory; nephrotropic; endocrine; viral infection; hepatitis B; hepatitis C; HIV; herpes simplex; cytomegalovirus; human papillomavirus; respiratory syncytial virus; influenza virus; restenceis; neurodegeneration; cancer; neurological; prion; inflammatory; autoimmune; pulmonary; renal; liver; mitochondrial; reproductive disease; chemical modification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid;
short interfering nucleic acid; siNA; expression; replication;
inhibition; RNA interference; virucide; anti-HIV; hepatotropic;
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                                                                                                  20-FEB-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADA13874 standard; RNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer used to isolate DNA encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 BP; 4
; 2002US-0358580P.
; 2002US-0363124P.
; 2002US-0386782P.
; 2002US-0406784P.
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                                                                                                  2003WO-US005346.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acid (siNA) oligonucleotide SEQ ID NO:211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ΒP
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Pred. No. 2
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of a target gene, where the siNA molecule comprises no ribonucleotides and each strand of the codube-stranded siNA comprises about 21 nucleotides. Also described: (1) a siNA molecule that inhibits replication of a virus and optionally does not require presence of a ribonucleotide for inhibition; (4) a siNA molecule that inhibits expression of a target gene and does not require presence of a ribonucleotide for inhibition; (4) a siNA molecule that inhibits expression of a target gene and does not require presence of a ribonucleotide for inhibition; (4) a siNA molecule that inhibits expression of a target gene by mediating RNA interference; and (5) a method for modulating expression of a gene in a cell using siNA complexity of an expression of a gene in a cell using siNA complexity of an expression of a target gene, inhibiting conversion of target RNA, and contininflammatory, plant antiviral, vasotropic, neuroprotective, cytostatic, cardiovascular, immunosuppressive, respiratory, mephrotropic expression of target genes, inhibiting expression of target RNA, and compression of target penes, inhibiting expression of target RNA, and content of any disorder that responds to modulation of gene expression, cespecially animal and plant viral infections, specifically hepatitis B or cytostally animal and plant virals infections, specifically hepatitis B or cytostally animal and plant virals infections, specifically hepatitis B or cytostally animal and plant virals, slNA molecules can be used: (a) for cytostally animal and plant virals infections, specifically hepatitis B or cytostally animal and plant virals. Infections, pocifically hepatitis B or cytostally animal and plant virals, slNA molecules such as createnesses such as createnesses and cytostally characters, binding affinity and/or mediates intoreases; and (b) for diagnosis, target can be used: (a) for diagnosis, target can be used: (a) for diagnosis,
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09-SEP-2002;
,15-JAN-2003;
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Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g.
                                     designed to target many related genes containing a conserved sequence. The present sequence represents a siNA oligonucleotide sequence, which used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 138;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-689785/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  short interfering nucleic acid containing no ribonucleotides, useful for treating viral infection, downregulates expression of target
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2002US-0409293P.
2003US-0440129P.
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Fosnaugh K,
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Mokler V, Jamison
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Sequence
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BP; 3 A; 7 C; 8 G; 0 T; 3 U; 0 Other;
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RESULT 3338
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Matches
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              human leukocyte antigen; HLA; probe; PCR; ss
                              Probe sequence
                                              04-DEC-2003
                                                                              ADB78519
                                                               ADB78519;
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                                                                                                                                221 CGGGACCCTCGGGAGCAGCTG 241
                                                                                                                       21
                                                                                                                                                       l Similarity
17; Conserv
                                                                              standard;
                                                                                                                       CGGGGCCCTCCAGAGCATCTG 1
                                                                                                                                                       Conservative
                                              (first entry)
                              #22 related to
                                                                               DNA;
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                                                                              21
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                                                                                                                                                                       Score 14.6;
                              the invention.
                                                                                                                                                       Mismatches
                                                                                                                                                               No.
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RESULT 3339
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to the detecting of a target nucleic acid sequence on a sample nucleic acid strand. The methods are useful for detecting the presence or absence of target nucleic acid sequences on sample nucleic acid strands that are characteristic of pathogens or gene variations and mutations relating to human leukocyte antigen (HLA) or T-cell receptor gene sequences, e.g. for HLA tissue typing, detecting genetically inherited diseases or detecting infectious organisms in tissues. The diagnostic probes are useful for detecting the presence of particular target nucleotide sequences. The present invention provides improved methods of detecting sample/target nucleic acid sequences, where the use of diagnostic probes having increased specificity reduces the number of alleles detected, which increases the resolution of the method, and does so at a lower cost. The present sequence represents a probe of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting the presence of a target nucleic acid sequence on a sample nucleic acid strand, useful for human leukocyte antigen tissue typing, comprises contacting a sample with a diagnostic probe under hybridizing
Saito K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 29; 62pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-363216/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-2001; 2001US-0324421P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-SEP-2002; 2002WO-US030238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003027309-A2
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                                                     24-SEP-2001;
                                                                                                                                   WO2003027309-A2
                                                                                                                                                              Synthetic
                                                                                                                                                                                         human leukocyte
                                                                                                                                                                                                                  Probe sequence #25 related to the invention.
                                                                                                                                                                                                                                                                                                ADB78522 standard; DNA;
                                                                               24-SEP-2002; 2002WO-US030238.
                                                                                                           03-APR-2003
                          (ONEL-) ONE LAMBDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention.
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                                                                                                                                                                                                                                                                                                                                                                    21
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17; Conser
                                                                                                                                                                                                                                                                                                                                                                                     GCCGTGCAGGTTCACCAGGAT 2760
                                                                                                                                                                                                                                                                                                                                                                     GCCGCGCAGGTCCCCCAGGTT 1
Lee J, Blair L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lee J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                    2001US-0324421P
                                                                                                                                                                                                                                             (first
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                                                                                                                                                                                         antigen; HLA; probe; PCR; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A.
                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                  0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8 C; 8
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                                                                                                                                                                                                                                                                                                 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                      .3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
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                                                                                                                                                          CC The present invention relates to the detecting of a target nucleic acid sequence on a sample nucleic acid strand. The methods are useful for CC detecting the presence or absence of target nucleic acid sequences on CC sample nucleic acid strands that are characteristic of pathogens or gene CC variations and mutations relating to human leukocyte antigen (HLA) or T-CC cell receptor gene sequences, e.g. for HLA tissue typing, detecting CC genetically inherited diseases or detecting infectious organisms in CC tissues. The diagnostic probes are useful for detecting the presence of particular target nucleotide sequences. The present invention provides comproved methods of detecting sample/target nucleic acid sequences, where CC the use of diagnostic probes having increased specificity reduces the number of alleles detected, which increases the resolution of the method, CC and does so at a lower cost. The present sequence represents a probe of the invention.
                                                                                             Query Match
Best Local 9
                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting the presence of a target nucleic acid sequence on a sample nucleic acid strand, useful for human leukocyte antigen tissue typing, comprises contacting a sample with a diagnostic probe under hybridizing
                                                                                                                                                        Sequence 21 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-363216/34.
                     2740 GCCGTGCAGGTTCACCAGGAT 2760
21 GCCGCGCAGGTTCCGCAGGCT 1
                                                                          l Similarity
17; Conserv
                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29; 62pp; English.
                                                                                                                                                          A; 8 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                               0.2%;
                                                                            0
                                                                                               Score 14.6;
Pred. No. 2
                                                                                Mismatches
                                                                                                 .3e+03
                                                                                                                    DB 1;
                                                                                                                    Length
                                                                                Indels
                                                                                                                      21;
                                                                                <u>.</u>
                                                                                Gape
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RESULT 3340
ADC16367
30-JUL-2001; 2001US-0308640P.
08-APR-2002; 2002US-0370970P.
                                                                                                                                                                                                                                                                                                                 ADC16367 standard; RNA;
                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                      proliferative
                                                                                                                                                                                                                                        expression interference; expression inhibition; target gene;
                                                                                                                                                                                                                                                          Short interfering
                                                                                                                                                                                                                                                                             18-DEC-2003
                                                                                                                                                                                                                                                                                                ADC16367;
                                   WPI; 2003-248169/24.
                                                     Caplen NJ,
Kallioniemi
                                                                                         (USSH )
(CARN-)
                                                                                                                                              30-JUL-2002; 2002WO-US024226
                                                                                                                                                                 13-FEB-2003
                                                                                                                                                                                  WO2003012052-A2
                                                                                (IMCO-)
                                                                                                                                                                                                                                interfering
                                                                              US DEPT HEALTH & HUMAN SERVICES. CARNEGIE INST WASHINGTON. IMPERIAL COLLEGE INNOVATIONS LTD
                                                   Morgan RA, Fire A, O, Cornelison JR,
                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                       disease; cancer; ds.
                                                                                                                                                                                                                               double stranded RNA; cytostatic; gene therapy;
                                                                                                                                                                                                                                                           double-stranded RNA oligonucleotide
                                                                                                                                                                                                                                                                                                                  21
                                                                                                                                                                                                                                                                                                                  ВP
                                                     Alton
                                                     Parrish S, Mousses S;
Alton EW, Griesenbach
                                                                                                                                                                                                                                                            SEQ
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                                                                                                                                                                                                                                                            ij
                                                                                                                                                                                                                                                             NO:92
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New RNA comprising double stranded RNA and a 3' or 5' overhang having a length of 0-nucleotide to 5-nucleotides on each strand, useful as reverse genetic and/or therapeutic tools for interfering or inhibiting expression

Claim

71; SEQ ID NO

92; 176pp;

English

of a target gene

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cc inhibition of expression of a target gene, where (I) comprises double ctranded RNA of 15-40 nucleotides in length and a 3' or 5' overhang chaving a length of 0-nucleotide to 5-nucleotides on each strand, where the sequence of the double stranded RNA is substantially identical to a gortion of a mRNA or transcript of the target gene. Also described: (I) cc interfering with or inhibiting the expression of a target gene in a cell cc yexposing the cell to an amount of (I); (2) a gene silencing array cc comprising a substantially flat substrate, and addressably arrayed cc different double-stranded RNA; (3) an array-based method of assessing a cc phenotypic effect of a double-stranded RNA on a target gene; (4) cc validating a gene as a potential drug target for a disease or condition; (5) selecting an optimised sequence of a double-stranded RNA for condition and (6) a short double-stranded RNA effective for interfering with or condition expression of a target gene in a cell; and (6) a short double-stranded RNA effective for interfering with or condition expression of a target gene comprising any of 311 20-78 cc conclectide sequences (see ADC16586). (I) has cytostatic confidence of a target gene therapy. The RNAs are useful as reverse genetic and/or therapeutic tools for interfering or inhibiting expression of a target gene. They are useful for treating proliferative diseases, confidence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                           Diagnosing c
patient, by
                                             cancer-associated
                                                                                                                                                                                                        D'andrea
                                                                                                                                                                                                                                                           (DAND )
(UYOR-)
                                                                                                                                                                                                                                                                                                                                    02-NOV-2001;
02-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC42515;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC42515 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention descrinhibition of expression of stranded RNA of 15-40 nucleous transfer in the stranded RNA of 1
                                                                         patient,
                                                                                                                                                       WPI; 2003-441436/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003039327-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chemosensitising;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer; Fanconi Anaemia; FA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FANCD2 PCR primer MG763 SEQ ID NO:181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2761 ACTCTGCGCACCATGTACTTC 2781
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                                                                                                                                                                                                                                                        DANA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21
                                                                                              or determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B₽;
                                             or determining cancer or increased risk of cancer in a
testing Fanconi Anemia/BRCA pathway gene or protein for a
diated defect, that indicates cancer or increased risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
ilarity 52.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes an RNA (I)
                                                                                                                                                                                                                                                        FARBER CANCER
OREGON HEALTH
                                                                                                                                                                                                                                                                                                                                 2001US-00998027.
2001WO-US045561.
                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US018153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                        Taniguchi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; 4
                                                                                                                                                                                                        'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; BRCA; cytostatic; microarray;
primer.
                                                                                                                                                                                                                                                           INST
SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
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                                                                                                                                                                                                        Timmers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.6; D
Pred. No. 2.3e
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ç;
                                                                                                                                                                                                        Grompe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .3e+03;
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                                                                                                                                                                                                        Fox
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The
                                                                                                method is useful for determining if a patient has cancer, or is at increased risk of developing cancer, e.g. breast, ovarian or prostate cancer. A microarray of the invention is useful for determining if a patient has cancer, or is at increased risk of developing cancer, by hybridising a nucleic acid sample to the nucleic acid sequences from the array, and detecting the presence of mutations in FA/BRCA pathway genes in the nucleic acid sample from the patient, where detecting the presence of mutations is indicative of a patient who has cancer, or is at increased risk of developing cancer. A method of the invention is useful for screening a chemosensitising agent, and the agent obtained is useful for screening a chemosensitising agent, and the agent obtained is useful
                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel method of diagnosing or determining if a patient has cancer or is at increased risk of cancer, involving testing a Fanconi Anaemia (FA)/BRCA pathway gene or protein for the presence of a cancer-associated defect, where the presence of one or more cancer-associated defects is indicative of cancer or an increased risk of cancer in the patient. The method of the invention has cytostatic activity. The
Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; SEQ ID NO 181; 160pp; English
                      treating a patient having a cancer. The present sequence is used in exemplification of the invention.
BP; 7 A;
7 C; 3 G; 4 T; 0
Ģ,
0 Other;
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Query
Best I
                                   Matches
                                            Local
                2742 CGTGCAGGTTCACCAGGATAC 2762
_
                                  l Similarity
17; Conser
CATTCAGATTCACCAGGACAC
                                  Conservative
                                          0.2%;
                                  0,
                                             Score 14.6;
Pred. No. 2
                                   Mismatches
                                             2.3e+03
                                                   DB 1;
                                                   Length
                                   Indels
                                  0
                                  Gaps
                                   0
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JP2002291485-A.
                                                                                                   Synthetic
                                                                                                          atopic dermatitis; steroid; RNaseA; RNase k6 precursor; PCR primer;
                                                                                                                 RNaseA PCR primer
                                                                                                                         18-DEC-2003
                                                                                                                                ADC64837;
                                                                                                                                       ADC64837 standard;
                                                                                                                        (first entry
                                                                                                                 SEQ ID NO: 2.
                                                                                                                                       DNA;
                                                                                                                                       21
                                                                                                                                       ВP
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88.

08-OCT-2002

03-APR-2001; 2001JP-00104621

03-APR-2001; 2001JP-00104621.

(KOKU-) (GENO-) GENOX SOYAKU KEN KOKURITSU SHONI KENKYUSHO O KK.

WPI; 2003-375782/36.

diseases. Examination of dermatitis for responsiveness to simproved treatment steroids in patients with atopic t of patients suffering from alle suffering from allergic

Example 2 Page 14; 24pp; Japanese.

The present invention describes a method for examining the response of patients with atopic dermatitis to steroids. The method comprises: (a) determining the expression levels of an RNaseA gene or an RNase k6 precursor gene in a living sample; and (b) comparing the response level with that of healthy subjects and steroid responsive patients, determined by PCR of CDNA. The present sequence represents an RNaseA PCR primer, which is used in an example from the present invention.

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Sequence 21 BP; 2 A; 5 C; 6 G; 8 T; 0 U; 0 Other;

밁

1 TGGAGTCTTTGCACGCCTCGG 21

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RESULT 3343
ACF36397
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                                                   Matches
                                                                Query Match
Best Local :
                                                                                                                                                          remaining nucleotides 5-17 are 2'-deoxy; and (b) the cytosines at positions 1, 4 and 19 are 5-methylated. Oligonucleotide shown in sequence ACP36398 (1) is used: (a) to delay progression of androgen-sensitive prostatic cancer cells to the androgen-independent state, in vivo or in vitro; (b) to treat prostatic cancer (after initially withdrawing androgens to induce apoptosis); and (c) to increase sensitivity of cancer cells (prostatic, renal, non-small cell lung, urothelial transitional, ovarian and some breast cancer cells) that express abnormal levels of TRPM-2 to chemotherapy or radiation. The modifications present in (1) increase stability in vivo and activity (both in vivo or in vitro) and result in a synergistic increase in effect when (1) is used with chemotherapeutic agents or other antisense oligonucleotides directed against other antispoptotic genes. The present sequence represents an anti-apoptotic protein TRPM-2 (testosterone-repressed prostate message-2)
                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a compound consisting of an oligonucleotide with a phosphorothicate backbone throughout, in which: (a) sugars on nucleotide residues 1-4 and 18-21 are 2'-O-methoxyethyl modified, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New modified antisense oligonucleotide, useful particularly for treating prostatic cancer, inhibits the testosterone-repressed prostate message-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gleave M, Rennie PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TRPM-2; testosterone-repressed prostate message-2; cytostatic; androgen; prostate cancer; anti-apoptotic protein; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TRPM-2
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                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Page 40; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-FEB-2002; 2002US-00080794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2003; 2003WO-US005305
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                               antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYBR-) UNIV BRITISH COLUMBIA
674 TGGAGTCTGTGCAAGCCCTGG 694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                   17;
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                                                                  Similarity
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                                                                                                                 21
                                                                                                               BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide.
                                                                                                               A;
                                                               0.2%;
                                                                                                               6 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Miyake H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
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                                                                                                               G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.6; DB 1;
Pred. No. 2.3e+03;
                                                                Score 14.6; DB 1; Length 21; Pred. No. 2.3e+03;
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                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nelson C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 21;
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                                                   Indels
                                                 0
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                                                 Gaps
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RESULT 3344
ADD14424/c
                          The present invention describes a predictor set comprising a plurality of colynucleotides or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein tyrosine kinase activity of cells, comprising whether a compound is capable of condulating the activity of cells, comprising whether a compound is capable of cells, comprising whether the cells express a plurality of markers, and correlating the expression of the markers to the compound's ability to correlate with compound sensitivity or resistance of cell lines for correlate with compound sensitivity or resistance of cells associated with a disease state, and (3) identifying polymucleotides and colypeptides whose expression pattern of a microarray of polymucleotides or polypeptides that predict compounds, analysing the expression pattern of a microarray of polymucleotides or polypeptides, and selecting polymucleotides or polypeptides that predict the sensitivity or resistance of cells associated with a disease state, comprising subjecting the plurality of colymucleotides or polypeptides that predict the sensitivity or resistance of cells associated with a disease state by using the expression pattern of a microarray. The polymucleotides and complete the polymucleotides or polypeptides and selecting the polymucleotides or polypeptides, and selecting the compounds that interact with protein tyrosine kinases and/or protein tyrosine kinase pathways. These may be used in determining drug sensitivity in patient response at a molecular level. The present of cancer) based on patient response at a molecular level. The present of cancer is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         predictor set; protein tyrosine kinase activity modulator;
protein tyrosine kinase pathway; protein tyrosine kinase; cytostatic;
gene therapy; drug sensitivity; genetic profile; cancer; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD14424 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotides and polypeptides for predicting the activity of compounds that interact with protein tyrosine kinases and/or protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JAN-2003; 2003WO-US001981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tyrosine kinase pathways.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Huang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-JAN-2002; 2002US-0350061P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            biomarker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fairchild CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 613; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shaw P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID
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Query Match Best Local Similarity

0.2%;

Score 14.6; DB 1; Pred. No. 2.3e+03;

DB 1;

Length 21;

Seguence

21

BP; 5 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

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RESULT 3345
ADC84417
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                                                                                                           RESULT 3346
ADD20200/c
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                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                              The invention comprises oligonucleotides for detecting and identifying subtypes of human papilloma virus (HPV) contained in a sample. The oligonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an oligonucleotide that was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                               Detector for identifying human papilloma virus subtypes, comprises carrier having two parts carrying first and second oligonucleotides that respectively hybridize with DNA contained in first and second subtypes of
single nucleotide polymorphism; SNP; fish; Salmo salar;
Orechromis niloticus; Atlantic halibut; microsatellite; cod;
                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-432398/41
                                                                                                                                                                                                                                                                                                                                                                                                                                    Heu
                                                                                                                                                                                                                                                                                                                                                                                                                                             Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-OCT-2001; 2001EP-00123379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EP1302550-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HPV detection method-related oligonucleotide Gap21-5
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                                 Oreochromis
                                                       15-JAN-2004
                                                                            ADD20200;
                                                                                                 ADD20200 standard; DNA;
                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probe; human papilloma virus; HPV; detection; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC84417;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KING-)
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Shih C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KING CAR FOOD IND CO LTD
                                                                                                                                                                   CCATCGCCTGCTTCGCACCCC 1582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTACTTTAGACCTAGCCCAGG
                                                                                                                                                      CCACCAACTGCTTAGCACCCC 21
                                                                                                                                                                                                                                          21 BP; 5 A; 11 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTCCCTTAGAGCTAGCCAAGG
                                                                                                                                                                                                                                                                                                                                   SEQ ID
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                                niloticus microsatellite primer
                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001EP-00123379
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                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 You C,
Yeh C,
                                                                                                                                                                                                                                                                                                                                 NO 647; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                          0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Huang H,
                                                                                                 21
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                                                                                                                                                                                                           Score 14.6;
Pred. No. 2
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                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 Pan C,
                                                                                                                                                                                                          DB 1;
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Chan
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                                 ID NO:835.
                                                                                                                                                                                                                     Length
                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Fan
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Novel isolated nucleic acid molecule comprising single nucleotide polymorphism associated with fish, useful for forming PCR primers which are used for detecting single nucleotide polymorphisms in fish nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polymorphic site; seabass; salmonidae; Tilapia; rainbow trout; halibut;
detection; primer; ss.
Claim 18; SEQ ID NO 835; 233pp; English.
                                                                                                                                                                                    WPI; 2003-627388/59
                                                                                                                                                                                                                                     Lie O,
                                                                                                                                                                                                                                                                                                                                        18-JAN-2002; 2002US-0349950P
16-AUG-2002; 2002US-0404200P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oreochromis niloticus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                           (GENO-)
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                                                                                                                                                                                                                                     Hoyum
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Sequence 21 BP; 9 A; 2 C; 8 G; 2 T; 0 U; 0 Other;

밁 ફ Matches Query Match 5318 CTCTCCTTTTCTCTCTTTGCC 5338 17; Conser Conservative 0.2%; 0, Score 14.6; Pred. No. 2 Mismatches .3e+03 DB 1; Length Indels 21; 0 Gaps 0;

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ADE16100/c
ID ADE11
XX
AC ADE10
ADE 29-JJ
DT 29-JJ
DX G-CO
XX
G-CO
XW G-CO
XW W antiji
XW antiji
XW angi
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ADD95120/c
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                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 17
                                                             G-coupled virucide;
                                                                                                                                                                                                                                                                                                                                                                 The invention relates to decreasing pigmentation in the skin, hair, wool or fur of a vertebrate or a mammal. This method comprises administering a composition comprising bone morphogenetic protein 4 (BMP-4), an active fusion protein of BMP-4, an active fragment of BMP-4, a BMP-4 mimic or a combination of any of those. The method is useful for lightening or decreasing pigmentation in the skin, hair, wool or fur of a vertebrate or a mammal. The current sequence represents a PCR primer for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Decreasing skin, hair, wool or fur pigmentation in a mammal comprising administering a composition comprising bone morphogenetic protein 4 (BMP-4), an active fusion protein or fragment of BMP-4, a BMP-4 mimic or its
                                  virucide; fungicide; cytostatic; nootropic; neuroprotective; antiparkinsonian; haemostatic; antilipaemic; neurogenesis; cell differentiation; cell proliferation; hematopoiesis; wou
                                                                                                G-coupled
                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                        amplification of BMP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003086313-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD95120 standard;
                        angiogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             combination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Decreasing skin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-APR-2002; 2002US-0372523P
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                                                                                                                                                                         ADE16100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-903051/82.
                                                                                                                          29-JAN-2004
                                                                                                                                                                                                                                                            1915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer;
                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                  1 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UNIV BOSTON.
                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                          AAACTTGGTGGCATTAACAAC 1935
                                                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Park
                                                                                                                                                                                                                                   AATGGTGGTGGCATTTACAAC 1
                                                             protein rec
fungicide;
                                                                                                protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID
                                                                                                                                                                                                                                                                                                                                  BP; 6
                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Å,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         type 1B upstream primer
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                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 skin; hair; wool; fur; bone morphogenetic protein 4; BMP-4;
                         gene therapy;
                                                                        receptor; antidiabetic;
                                                                                                receptor related forward
                                                                                                                                                                                                                                                                                                                                  P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NO 4; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Botchkarev
                                                                                                                                                                        DNA;
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                                                                                                                                                                                                                                                                                             0.2%;
                                                                                                                                                                                                                                                                                                                                  6 C; 3
                                                                                                                                                                                                                                                                                                                                                        receptor type 18 DNA.
            pharmacogenomics;
                                                                                                                                                                         21 BP
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              chromosome mapping; ti
cogenomics; human; PCR;
                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                  G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                              Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                  Mismatches
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                                                                        anorectic;
                                                                                                                                                                                                                                                                                                         DB 1;
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                                                                                                primer,
                        tissue typing;
                                                                                                                                                                                                                                                                                                         Length 21;
                                                                                                                                                                                                                                                                                    Indels
                                                                           antibacterial;
                                                                                                SEQ
                                      wound healing;
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The invention relates to a novel isolated G-coupled protein receptor cc related polypeptides. The novel polypeptide comprise any of the 22 fully defined sequences of 87-1780 amino acids, given in the specification; their mature forms; and possible variants. The novel polypeptides have the following activities: antidiabetic, anorectic, antibacterial, controlled, fungicide, cytostatic, nootropic, neuroprotective, including a preventing in a human, a pathology associated with the G-coupled protein receptor related polypeptides are useful in a method of treating or converting in a human, a pathology associated with the G-coupled protein receptor related polypeptides. The polypeptides are useful in the complete receptor related polypeptides. The polypeptides are useful in the complete protein and follows a method of treating or complete related polypeptides. The polypeptides are useful in the complete protein and follows a method of treating, preventing or disposaing diseases, such as metabolic disorders, diabetes, obesity, infectious diseases, cancer-associated diseases, neurodegenerative disorders, and various dyslipidaemias; metabolic disturbances associated with obesity, metabolic X syndrome and wasting disorders associated with obesity, metabolic X syndrome and wasting disorders associated with chronic diseases and various cancers. The nucleic acids and polypeptides may also be used as targets for the identification of small molecules that modulate or inhibit e.g. neurogenesis, cell differentiation, cell proliferation, hematopoiesis, wound healing and angiogenesis, in gene content of the province of the province in the respective of disposes; to Novx substances for use in the respective of disposes; in chromospecifically to Novx substances for use in the respective of disposes; in chromospecifically to Novx substances for use in the respective of disposes; in chromospecifically to Novx substances for use in the respective of disposes; in chromospecifically to Novx substances for use in the respective of dispose
              13 APR-2001;
17-APR-2001;
19-APR-2001;
20 APR-2001;
23-APR-2001;
23-APR-2001;
24 APR-2001;
27-APR-2001;
30-MAY-2001;
30-MAY-2001;
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05-APR-2001;
05-APR-2001;
10-APR-2001;
13-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Voss EZ,
Zhong M,
Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated NOVX polypeptides and preventing, diagnosing or treating diabetes, obssity, dyslipidemias, diabetes, obssity, dyslipidemias,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pena
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31-JUL-2001;
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13-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alzheimer's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-067574/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gerlach V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rlach V, Liu X, Miller CE, Spytek KA, Zerhusen B
Shenoy SG, Zhong H, Smithson G, Casman SV, Bol
Vernet CAM, Macdougall JR, Rastelli L, Anderson
Mezes PD, Furtak K, Patturajan M, Burgess CE, M
RA, Taupier RJ, Edinger SR, Mazur A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; 2001US-0324775P.
; 2001US-0333900P.
; 2002US-00115479.
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2001US-0286068P.
2001US-02872119.
2001US-0288509P.
2001US-0294801P.
2001US-0294801P.
2001US-0394801P.
2001US-0394801P.
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2001US-0285325P.
2001US-0285609P.
2001US-0285748P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 130; 320pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001US-0283678P
2001US-0283687P
2001US-0283710P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer, Parkinson's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotides, useful for NOVX-associated disorders e
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Zerhusen BD; an SJ, Boldog

FL;

Malyankar UM;

e.g.

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RESULT 3350
AAQ32173/c
ID AAQ32173 E
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AAQ14243
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                                                                                      Query Match
Best Local S
Matches 17
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Best Local S
Matches 17
                                                                                                                                      The primer is used with primer CMLII (AAQ14244) to amplify chimeric mRNA contg. a specific exon-exon junction associated with chronic myeloid leukaemia (CML). It is complementary to CDNA sequences within the break point cluster region (BCR) exon 2. It can distinguish between ALL BCR-ABL chimeric mRNA and CML BCR-ABL mRNA. The CML DNA sequences used to design the primer are reported by Heisterkamp, N., et al. Nature 315:758 (1985); Grosvelt, G., at al Mol. Cell Biol. 6:607 (1987); and Shtivelman, E., et al Cell 47:277 (1986). See also AAQ14241-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tissue typing, preventive medicine, and pharmacogenomics. polynucleotide sequence represents a primer relating to tl coupled protein receptor related polypeptides of the investigation.
                                                                                                                         Sequence
                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                         Method for detecting chimeric mRNA - useful e.g. for distinguishing between acute lymphocytic leukaemia and chronic myeloid leukaemia.
                                                                                                                                                                                                                                                                     WPI; 1991-324515/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ14243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ14243 standard; DNA;
                                                                                                                                                                                                                                                                                                                         05-AUG-1988;
                                                                                                                                                                                                                                                                                                                                         05-AUG-1988;
                                                                                                                                                                                                                                                                                                                                                           15-OCT-1991.
                                                                                                                                                                                                                                                                                                                                                                             US5057410-A
                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                              Acute lymphocytic leukaemia; chimeric; mRNA; ABL; breakpoint; cluster; BCR; ALL; exon junction; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer CMLI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17; Conserv
                                                     N
                                                                                      l Similarity
        standard;
                                                                 GAGCTCCAGATCCTGACGTAC 2559
                                                                                                                                                                                                                                                                                      ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAAGACAGCAAGGAGGAGTGA 1
                                                                                                                       22 BP; 6
                                                                                                                                                                                                                       Page 13; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGAGGAAGCAAGGAGGAGGA 2879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 BP; 1 A;
                                                     GAGCTGCAGATGCTGACCAAC
                                                                                     0.2%;
nilarity 81.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
ilarity 81.0%;
Conservative
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                                                                                                                                                                                                                                                                                      Mccormick
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                                                                                                                        A; 6 C; 7 G; 3 T; 0 U; 0 Other;
         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 C; 2 G; 10 T; 0 U; 0 Other;
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         22
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                                                                                     ; Score 14.6; D
; Pred. No. 2.4e
0; Mismatches
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Pred. No. 2.3e+03;
0; Mismatches 4
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                                                                                              .4e+03;
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                                                                                                        DB 1;
                                                                                                      Length 22;
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01-MAY-1992; 12-NOV-1992. WO9219739-A1

92WO-US003624

Bacillus thuringiensis

crops; nematode Reverse

pests;

worms; nemacricine pests; CryV proteins

PCR

primer for novel nematode active genes eg BT toxins.

nematicidal

toxin; agriculture;

plants;

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RESULT 3351
AAQ30933/c
                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                             Query Match
Best Local
              27-AUG-2003
25-MAR-2003
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31-JAN-1992;
23-APR-1992;
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25-MAR-2003
30-MAR-1993
                                                                                                                                                                                                                                                                                                This
of a
2003
                                                                                                                                                                                                                                                                                                                                                                            Nematocidal toxins from Bacillus thuringiensis animal or plant parasites, and DNA acid coding hosts and transgenic plants.
  30-MAR-1993
                                                       AAQ30933;
                                                                                  AAQ30933
                                                                                                                                                                                                                                                                    Sequence 22
                                                                                                                                                                                                                                                                                                                                                     Claim 1(f);
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MYCO ) MYCOGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-DEC-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP517367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacillus thuringiensis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nematode worms; nematicide; crops; pests; CryV proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ32173;
                                                                                                                                                      4405 TITACAAAAATGAATTITTCC 4425
|:|:::|| ||||| ||||
22 TWGAYMRAATTGAATTATTCC 2
                                                                                                                                                                                                                                                                                             degenerate sequence represents novel nematicdal toxin from BT to correct PN field.) (Updated
                                                                                                                                                                                                               14;
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                                                                                                                                                                                                                          0.2%;
Similarity 66.7%;
                                                                                    standard;
                                                                                                                                                                                                                                                                    BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                     Page 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer for novel nematode active genes eg BT toxins.
(revised)
(revised)
(first entry)
                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Schwab
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(revised)
(first en
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92US-00830050.
92US-00871510.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nematicide; nematicidal toxin; agriculture;
                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                    2 C; 2 G; 7 T; 0 U; 4 Other;
                                                                                                                                                                                                                                                                                                                                                     57pp; English.
                                                                                    22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Payne
                                                                                    BP
                                                                                                                                                                                                             4.
                                                                                                                                                                                                                            Score 14.6; DB 1;
Pred. No. 2.4e+03;
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                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                              a reverse PCR primer for the cloning as in AAQ32172. (Updated on 25-MAR-on 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Narva
                                                                                                                                                                                                                                         DB 1;
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    useful for control of
sequences, transformed

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                                                                                                                                                                                                             Indels
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RESULT 3352
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Best Local
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31-JAN-1992;
23-APR-1992;
AAQ90299 and AAQ90300 are a pair of upstream and AAQ90301 and AAQ90302 are a pair of downstream primers for the PCR amplification of AAQ73001 and AAQ73002, which encode AAA62657 and AAA62658 human tissue plastin plastin) and human leukocyte plastin (1-plastin) respectively. The plastin cDNA and amino acid sequences were used to develop isoform specific plastin nucleotide probes, and isoform specific anti-plastin
                                                                                                                                                                                                                                                                                                  nucleotide
                                                                                                                                                                                                                                                                                                            Human plastin;
                                                                                                                                                                                                                                                                                                                               Human plastin
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11-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                AAQ90300;
                                                                                                                                                                                                                                                                                                                                                                                                   AAQ90300 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1(f);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New genes and Thuringiensis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1992-398866/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MYCO)
                                                                                                                                                                                        07-JUN-1988;
16-MAR-1990;
                                                                                                                                                                                                                                          01-NOV-1994.
                                                                                               DNA encoding leukocyte-plastin and tissue-plastin - prode for distinguishing human haemopoietic cells,
                                                                                                                                                  Lin
                                                                                                                                                                                                                      10-JAN-1991;
                                                                     Example 2; Col 19; 17pp; English.
                                                                                         and neoplastic cells.
                                                                                                                               WPI; 1994-349444/43.
                                                                                                                                                                      (CALY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4405
                                                                                                                                                  ú
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  degenerate sequence represents novel nematicdal toxin from BT to correct PN field.) (Updated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 66.
                                                                                                                                                  Aebersold RH,
                                                                                                                                                                      CALIFORNIA INST OF TECHN.
                                                                                                                                                                                                                                                                                                                                                                                                                                              TWGAYMRAATTGAATTATTCC 2
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                                                                                                                                                                                                                                                                                                  in; haemopoietic cells; neoplastic; PCR probes; anti-plastin antibodies; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schwab GE,
                                                                                                                                                                                                                                                                                                                                                   (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            toxins against nematodes - obtd. fr isolates with nematocidal activity.
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92US-00830050.
92US-00871510.
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                                                                                                                                                                                                                                                                                                                               PCR primer.
                                                                                                                                                                                                                                                                                                                                                   entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 epresents a reverse PCR primer for
n from BT as in AAQ30943. (Updated
(Updated on 27-AUG-2003 to correct
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Pred. No. 2.4e
4; Mismatches
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                                                                                                   used to develonormal tissue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    correct OS field.)
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RESULT 3353
AAQ61522/c
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                                       The Ikaros gene encodes a zinc finger protein which can be used in a CC therapeutic composition to treat animals with an immune system disorder. CC It may also be used for assessing whether a subject is at risk for an CC immune disorder. It is of particular use in treating a disorder of the CC corpus striatum. Heterologous genes may be expressed by placing them CC under the control of an Ikaros protein. Potential high affinity binding sites CC for the Ikaros proteins were found in the enhancer and promoter regions CC for the TR-alpha, 'beta and 'delta, the CD3-delta, -epsilon and -gamma CC genes, the SL3 and HIV long terminal repeat and in the regulatory domains CC motif were also found in the purine boxes of the Ikaros found in the purine boxes of the Ikaros for the TR the TDT promoter as well as in the NFKB variant sites of the CC LYF site of the TDT promoter as well as in the NFKB variant sites of the CC 2003 to add missing OS field.) (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
10-MAR-2003
21-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antibodies. Using the fact that human cells that express only 1-plastin are haemopoetic cells, and human cells that express both 1-plastin and t plastin are neoplastic, the above probes and antibodies could be used to distinguish between the above cell types. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 1 A; 18 C; 1
                                                                                                                                                                                                                                                                                                                               I-cell pathway regulatory gene, Ikaros - encodes family of unique finger proteins, useful for treating immune system disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  corpus striatum; regulatory gene; enhancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ61522 standard; cDNA;
                               field.)
                                                                                                                                                                                                                                                                                                 Disclosure; Page 28; 112pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             Georgopoulos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-MAR-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ikaros; zinc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ61522;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    finger; protein; immune disorder; therapy; treatme
tum; regulatory gene; enhancer; regulatory element;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enhancer element comprising Ikaros binding
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(revised)
(first en
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Pred. No. 2
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RESULT 3355
AAQ86627
ID AAQ8662
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AC AAQ8662
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AAQ86634
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                                                                                                                   Query Match
Best Local S
Matches 17
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Best Local S
Matches 17
                                                                                                                                                                        The oligonucleotide AAQ86634 is a non-promoter primer for the CML major breakpoint amplification region. It is used to illustrate that small changes in the NA sequence result in large changes in the amplification efficiency .AAQ86634 is capable of serving as a primer for the synthesis of autocatalytic oligonucleotides which require no change in the PCR conditions i.e.constant temperature, pH and ionic strength. This sequence is useful in generating multiple copies of specific nucleic acid target
         25-MAR-2003
15-NOV-1995
                                                                                                                                                                                                                                       Example 18; Col 47; 58pp; English.
                                                                                                                                                                                                                                                      Amplification of nucleic acid targets - using a reverse transcriptase with RNase H activity and a RNA polymerase at constant temp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
16-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ86634 standard;
                                AAQ86627;
                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                              WPI; 1995-130686/17.
                                                                                                                                                                                                                                                                                                                             11-JUL-1989;
10-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                    19-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                    21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                   US5399491-A.
                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                  Primer; autocatalytic;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Non-promoter primer for the CML major breakpoint region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ86634;
                                               AAQ86627
                                                                                                                                                                                                                                                                                                             (GENP-) GEN-PROBE INC.
                                                                                              2539 GAGCTCCAGATCCTGACGTAC 2559
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17; Conserv
                                                                                                                     1 Similarity
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                                              standard;
                                                                                      GAGCTGCAGATGCTGACCAAC
                                                                                                                                                                                                                                                                                              Kacian DL;
                                                                                                                                                                  (Updated on 25-MAR-2003 to correct PF
                                                                                                                                                  BP; 6 A;
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        (revised)
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(first entry)
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90US-00550837.
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                                              DNA;
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        entry)
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                                                                                                                          0.2%;
                                                                                                                                                  6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; target;
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Pred. No. 2.4e+03;
                                                                                                                           Score 14.6;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                  sequence; ss
                                                                                                                           .4e+03
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Best Local Similarity Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ86626-28 are primers and a probe for the CML chromosomal translocation. They are used to produce autocatalytic oligonucleotides which require no change in the experimental conditions i.e.constant temperature, pH and ionic strength. These sequences are useful in generating multiple copies of specific nucleic acid target sequences. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                   CML-2 chromosomal translocation major breakpoint; t(9; auto-catalytic; synthesis; RNA target sequence; assay;
                                                                                                                                                                                                                                                                                                 CML-2 chromosomal translocation major breakpoint t(9;22) (+) primer
                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
17-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                            AAT15572;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT15572 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amplification of nucleic acid targets - using a reverse transcriptase with RNase H activity and a RNA polymerase at constant temp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CML chromosomal translocation plus strand primer.
11-JUL-1989;
                                        10-JUL-1990;
                                                                                    02-JAN-1996.
                                                                                                                            US5480784-A.
                                                                                                                                                                                                               quantification;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUL-1989;
10-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2539 GAGCTCCAGATCCTGACGTAC 2559
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kacian DL;
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                                                                                                                                                                                                                                                                                                                                           (revised)
(first en
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90US-00550837
89US-00379501
                                          90US-00550837.
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                                                                                                                                                                                                                                                                                                                                           entry)
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Pred. No. 2.
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                                                                                                                                                                                                                                   22); primer;
detection;
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fultz TJ,

Kacian DL;

(GENP-)

GEN-PROBE

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RESULT 3357
AAT42418
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is a primer for the CML-2 chromosomal translocation major breakpoint t(9;22), which was used to demonstrate an improved method for synthesising multiple copies of a RNA target sequence. The method comprises combining the target with a primer which hybridises to the 3'-terminal portion of the target, a promoter primer which hybridises with a portion of the DNA primer extension prod., reverse transcriptase, RNase H and transcriptase. It can be used as a component of an assay to detect and/or quantitate specific target sequences in clinical, environmental or forensic samples. It also has the advantages of being autocatalytic, using the cooperative action of a DNA polymerase, e.g. reverse transcriptase and avoids repetitive manipulations of reaction conditions, e.g. temp., ionic strength and pH. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Auto-catalytic synthesis of multiple copies of an RNA target sequence - uses cooperative action of a DNA and RNA polymerase in presence of RNase H, useful for detection of target sequence e.g. in clinical or environmental sample.
                                                                                                                                                                                                                                                                                                                                                                                    HIV; probe;
BCL-2; PCR;
AAT42417-T42419 represent primers and a probe translocation \mathsf{t}(9;22). These sequences can be of the kits of the invention. The kits of the
                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                  CML chromosomal translocation primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
28-APR-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-068248/07
                                                                                        Detection of HIV nucleic acids in samples - using new specific oligo-
nucleotide(s) for the amplification and detection of target sequences
                                                                                                                                                                                                                                                                                                 11-SEP-1996
                                                                                                                                                                                                                                                                                                                              EP731175-A2
                                                                                                                                   WPI; 1996-403995/41
                                                                                                                                                                                                                         11-JUL-1989;
10-JUL-1990;
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                                                                                                                                                                                              (GENP-) GEN-PROBE INC
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAGCTGCAGATGCTGACCAAC 22
                                                                                                                                                                S
                                                          Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 6 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                     primer; amplify; polymerase chain reaction; microorganism;
hepatitis B virus; HBV; CML; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                                                           89US-00379501.
90EP-00307503.
                                                                                                                                                                                                                                                                      96EP-00101621
                                                          8; 66pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                           English.
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Pred. No. 2
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 for the CML chromosomal used in modified versions invention, are for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
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RESULT 3358
AAX83079
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detecting the presence of HIV nucleic acid sequences in a sample. The kits comprise two amplification primers (such as AAT40182), and a probe (such as AAT42404) for detection of the amplified sequence. By using these sequences, the amplification of HIV nucleic acid sequences is improved. The kits can also be used for the detection of other microorganisms, by using different probe sequences. Other sequences that can be detected using this method include those from HBV (using the sequences shown in AAT42410-T42412), and BCL-2 (using AAT42413-T42416). The samples can be clinical, environmental or forensic samples, and the method produces large amounts of the target sequence for a variety of uses. The method can also be used to produce multiple copies of a target sequence for use in cloning, and sequencing, and to produce probes for the target sequence. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Sim
Matches 17;
                                                                                                                                                                                                                                             29-DEC-1995;
29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
12-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 6 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                               Isolated nucleic acid molecule encoding the WRN gene product - useful
detection and treatment of Werner's syndrome, and related diseases.
                                                                                                                                                                                              Oshima
                                                                                                                                                                                                                                                                                                                            30-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                      10-JUL-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; WRN; Werner's syndrome; detection; diagnosis; autosomal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer 5ES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX83079;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX83079 standard;
                                                                                                                                                                    WPI; 1997-363671/33.
                                                                                                                                                                                                                                                                                                                                                                              WO9724435-A1
                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                            recessive disorder;
                                                                                                                                                                                                                        (DARW-) DARWIN MOLECULAR CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2539 GAGCTCCAGATCCTGACGTAC 2559
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                             phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
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                                                                                                                                                                                              Mulligan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; PCR; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in the human WRN gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                              Schellenberg GD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            mutation;
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Primers AAX83071-X83082 were used to PCR amplify, detect and identify mutations in the human WRN gene (AAX83003) which encodes a protein related to Werner's syndrome. The products can be used for the detection and treatment of Werner's syndrome (WS), an autosomal recessive disorder

Example

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48; 153pp; English.

Sequence with a

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complex phenotype,

as well as

related diseases

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RESULT 3359
AAX83072
ID AAX83077
XX AAX8307
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AC AAX8307
XX January
AC AAX8307
XX January
AC AAX8307
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XX Human,
KW recessi
XX Primer
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XX W097244
XX W097244
XX January
PN W097244
XX W097244
XX W097244
XX January
PN Jo-JUL-
XX January
PR 30-JAN-
PR 
  RESULT 3360
AAT68896
ID AAT6889
XX
AC AAT6889
XX
DT 06-APR-
XX
                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-DEC-1995;
29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
12-APR-1996;
                        06-APR-1998
                                                            AAT68896;
                                                                                                                                                                                                                                                                                                                                                                                    mutations in the human WRN gene (AAX03003) which encode related to Werner's syndrome. The products can be used and treatment of Werner's syndrome (WS), an autosomal r with a complex phenotype, as well as related diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                              Primers AAX83071-X83082 were used to PCR amplify, detect and identify mutations in the human WRN gene (AAX83003) which encodes a protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated nucleic acid molecule encoding the WRN gene product - useful detection and treatment of Werner's syndrome, and related diseases.
                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                recessive disorder; phenotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; WRN; Werner's syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX83072;
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                                                                                                                                                                                                                                    7337
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                                                                                                standard;
                                                                                                                                                                                                                                AGCTGTACCTTGTCCAGTCCA 7357
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                                                                                                                                                                                              AGATGTACTTTGGCCATTCCA 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 48; 153pp; English.
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                                                                                                                                                                                                                                                                      .larity 81.0%;
Conservative
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                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mulligan
                                                                                                  ВP
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                                                                                                                                                                                                                                                                                      Score 14.6;
Pred. No. 2.
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detection; diagnosis; autosomal; mutation;
primer; PCR; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in the human WRN gene
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                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                  U; 0 Other;
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                                                                                                                                                                                                                                                                                                              DB 1;
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                                                                                                                                                                                                                                                                                                          Length 22;
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                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 3361
AAT91777
                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
Synthetic
                                  PCR; primer; amplify; polymerase chain reaction; chronic myelogenous leukaemia; human; bcr2-abl2; cytogenetic remission; Ph chromosome; bcr3-abl2; acute lymphotrophic leukaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This RT-PCR 5' primer is designed to a human breakpoint cluster region (BCR) and is used in a novel method of preparing drug-resistant, non-malignant haematopoietic cells. This method involves the construction of a new expression cassette comprising a first nucleic acid molecule which encodes resistance of a host cell to a cytotoxic agent, operably linked to a first promoter which functions in the host cell and a second nucleic acid molecule operably linked to a second promoter which functions in the host cell. The second nucleic acid molecule operably linked to a second promoter which functions in the host cell. The second nucleic acid molecule on a nucleic acid molecule or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide whose expression decreases the expression of an RNA or a polypeptide present in a malignant cell only. This method can eliminate residual neoplastic disease in a patient, where the disease has an immature hematopoietic progenitor cell with a well-defined gene rearrangement. Diseases such as chronic myelogenous leukaemia which is associated with a BCR/ABL gene rearrangement, acute lymphoblastic leukaemia and acute promyelocytic leukaemia may be treated using this
                                                                                                                                  Primer BB300b for bcr2-abl2 and bcr3-abl2 translocation regions
                                                                                                                                                                        25-MAR-2003
08-JAN-1998
                                                                                                                                                                                                                                AAT91777;
                                                                                                                                                                                                                                                                    AAT91777 standard; DNA; 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Expression cassette for forming drug resistant hematopoietic stem cells decreases RNA or protein found only in malignant cells; for treating leukaemia(s), such as chronic myelogenous leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-289281/26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                          2539
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                        GAGCTCCAGATCCTGACGTAC 2559
                                                                                                                                                                                                                                                                                                                                                                    GAGCTGCAGATGCTGACCAAC 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                          amplify; polymerase chain reaction; haematopoietic cell;
                                                                                                                                                                      (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RT-PCR
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Pred. No. 2.4e+03;
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Length Indels

22;

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Gaps

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translocation region; CML cell;

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ARV29432
ID AAV2943
XX AAV2943
AC AAV2943
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC AAT91749-T91763, and AAT91765-T91792 are primers used in the method of CC the invention. AAT91754-T91759 can also be used as capture CC also be used as detector agents. The method of the invention is for CC detecting or monitoring chronic myelogenous leukaemia (CML) cells in a CC human patient. The method comprises obtaining RNA from haematopoietic CC cells of the patient, and amplifying it using a pair of primers that CC amplify both the bcr2-abl2 and bcr3-abl2 translocation regions. The CC amplified sequence is contacted with a capture agent comprising a capture CC amplified for the bcr2-abl2 and bcr3-abl2 translocation regions. The CC mixture is contacted with a capture mixture. The capture ON is CC specific for the bcr2-abl2 and bcr3-abl2 translocation regions. The CC mixture is contacted with a solid phase is washed, then contacted with a CC detector agent comprising a detector ON specific for the bcr2-abl2 or bcr3-abl2 translocation regions and a label. The amount of labelled CC detector ON bound to the solid phase is then correlated with the presence CC or quantity of CML cells in the patients. It can also be used prognostically to assess cytogenetic remission in patients with CML. The method detector contacted with contacted with the bcr3-abl2 translocations associated with CML. The method detect CML in the absence of the ph chromosome, nor CC does it detect acute lymphotrophic leukaemia (ALL) even if the ALL cc field)
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 abl2 trans-location regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detection of chronic myelogenous leukaemia cells - RNA from haematopoietic cells with primers for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-AUG-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 10; 79pp; English
                                                                        Calcium ion channel alpha1 subunit; human; episodic ataxia type 2; familial hemiplegic migraine; FHM; EA-2; treatment; diagnosis;
                                                                                                                               Calcium ion channel alphal subunit exon 28
                                                                                                                                                                                                                                           AAV29432 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DADE-) DADE INT INC
   Homo sapiens
                                                                                                                                                                   31-JUL-1998
                                                         primer;
                                                                                                                                                                                                                                                                                                                                                                       2539 GAGCTCCAGATCCTGACGTAC 2559
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Similarity 81.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.6; DB 1; Length 22; Pred. No. 2.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                               specific reverse primer.
                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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10-SEP-1996; 02-OCT-1996; 30-OCT-1996;

96US-0025719P. 96US-0028056P. 96US-0029679P.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This primer is used for the PCR amplification of an exon of the human CC calcium ion channel alpha 1 subunit. The channel is related to familial CC hemiplegic migraine (FHM) and/or episodic ataxia type 2 (EA-2) and is CC derived from, related to or associated with a gene present in humans on CC chromosome 19p13.1-13.2. The encoding nucleic acid can be used to CC localise or identify genes related to episodic neurological disorders, CC specifically migraine, FHM or EA-2, but also epilepsy. It can also be CC used to distinguish between alleles of the corresponding gene. Cells and CC animals containing recombinant expression vectors comprising the nucleic acid can be useful in study, development and treatment of migraine, FHM, CEA-2 and epilepsy. Proteins or peptides encoded by the nucleic acid and CC diagnose FHM, EA-2, migraine and other neurological conditions associated CC with cation channel disfunction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 17;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human nucleic acid associated with migraine and episodic ataxia type 2 - useful for diagnosis and development of, e.g. familial hemiplegic migraine and episodic ataxia type 2.
                                                                                                                                                                                                                       Hepatocyte nuclear factor 4 alpha; HNF-4 alpha; MODY1; human; transcription factor; maturity onset diabetes of the young; I diabetes; NIDDM; diagnosis; therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22
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                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                      Hepatocyte nuclear factor 4
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                                                                                                                                                                              sapiens.
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Pred. No. 2.4e+03;
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                                                                                                                                   New nucleic acid encoding Ikaros protein involved in differentiation of lymphocytes - existing in several related products, used to treat e.g. immune diseases control cell differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bell GI,
Horikawa
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                                                                                                                                                                                                                                                                                  WPI; 1998-378292/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mIK; transcription factor; mouse; lymphocyte; ferentiation; T cell; cancer; immunodeficienc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell
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                                                                                                                                                                                                                                                                                                                                                                                                HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease;
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This oligonucleotide from the T cell receptor delta enhancer was identified as a potential high affinity binding site for Ikaros \cdot

proteins

AAV66352-55 represent CML-2 chromosomal translocation t(14;18) major breakpoint amplification region non-promoter primers. The primers are used to exemplify the invention. The specification describes methods o synthesising multiple copies of a target nucleic acid sequence autocatalytically under conditions of substantially constant temparatu ionic strength and pH are provided in which multiple RNA copies of the target sequence autocatalytically generate additional copies. The targe sequence is a block splice template which comprises two nucleic acid regions. The first region is located 3' of the second region and is blocked at its 3' terminus to inhibit primer extension to the conditional copies.

temparature,

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Example 18; Col 43; 51pp; English.

Disclosure;

Page

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158pp; English

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Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (see AAW70963-71). It includes the core motif GGGAA found in consensus recognition sequences for murine Ikaros protein isoforms mIk-1, mIk-2 and imIk-3 (see AAV52830-32). High affinity binding sites for Ikaros have been found in enhancer and promoter regions of the regulatory domains of the TCR antigen complex, the CD3 genes, the SL3 and HIV long terminal repeat and in the regulatory domains of other T cell restricted antigens (see AAV45358-402) by gel retardation assay. Ikaros is involved in early differentiation of lymphocytes. The invention provides Ikaros mucleic acids (see AAV42805-11 and AAV42840) and polypeptides, vectors and host cells. These are used to treat T and B cell diseases, to control expression of heterologous genes placed under control of an Ikaros responsive element, to treat nervous system diseases and to modulate cell division, amplification or differentiation, especially in haematopoietic cells.
                                                                                                                                                                             Block splice template useful for amplification of nucleic acids - comprises two nucleic acid regions, the first region located 3' of the second region and blocked at its 3' terminus to inhibit primer extension
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CML-2
block
                                                                                                                                                                                                                                   WPI; 1998-582557/49.
                                                                                                                                                                                                                                                             Fultz TJ,
                                                                                                                                                                                                                                                                                                               11-JUL-1989;
10-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                       06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                           US5824518-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CML-2
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                                                                                                                                                                                                                                                                                     (GENP-) GEN-PROBE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV66352 standard; DNA;
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                                                                                                                                                                    DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosomal translocation t(14; 18) major breakpoint; splice template; autocatalytic RNA amplification; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosomal
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                                                                                                                                                                   polymerase.
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                                                                                                                                                                                                                                                          Kacian DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 12
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90US-00550837.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  translocation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  major breakpoint non-promoter
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RESULT 3366
AAV66350
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Best Local (
Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                        Query Match
Best Local (
                                                                                                                                                                                                      AAV66349-50 represent CML-2 chromosomal translocation t(9;22) primers, for the (+) and (-) strands respectively. The primers are used to exemplify the invention, together with probe AAV66351. The specification describes methods of synthesising multiple copies of a target nucleic acid sequence autocatalytically under conditions of substantially constant temparature, ionic strength and pH are provided in which multiple RNA copies of the target sequence autocatalytically generate additional copies. The target sequence is a block splice template which comprises two nucleic acid regions. The first region is located 3' of the second region and is blocked at its 3' terminus to inhibit primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Block splice template useful for amplification of nucleic acids - comprises two nucleic acid regions, the first region located 3' of the second region and blocked at its 3' terminus to inhibit primer extension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22 BP; 6 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     recognised by acids, especia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-582557/49.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUL-1989;
10-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CML-2 chromosomal translocation t(9; 22); be autocatalytic RNA amplification; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CML-2 chromosomal translocation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymerase,
                                                                                                                                       extension by a DNA polymerase, and the second region comprises a promoter sequence recognised by an RNA polymerase. The methods are used to amplify nucleic acids, especially RNA, for analysis, cloning or probe production
                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENP-) GEN-PROBE INC.
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nes 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rase, and the second region comprises a promoter sequence ised by an RNA polymerase. The methods are used to amplify especially RNA, for analysis, cloning or probe production
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                                                                                                                                    acids, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymerase.
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                                                                                             22 BP; 6 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Col 9; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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90US-00550837.
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                      81.0%;
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Pred. No. 2.4e+03;
                        Score 14.6; DB 1
Pred. No. 2.4e+03
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  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22); block splice template;
                                                 DB 1;
  4
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                                              Length
    Indels
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  Gaps
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2539 GAGCTCCAGATCCTGACGTAC 2559

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14-JAN-1999 AAV67083;

(first entry)

AAV67083 standard; cDNA; 22

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Mouse TCR-delta enhancer deltaE5

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RESULT 3368
AAV67083/c
                                                                                                            Matches
                                                                                                                        Query Match
Best Local (
                                                                                                                                                                                                  PCR primers (see AAV45541 and AAV45542) are designed for the PCR amplification of the Helicobacter pylori babB gene. The invention relates to recombinant DNA (see AAV62460-61) comprising a fusion gene prepared by ligating an antigenic determinant coding gene (e.g. the babB gene) of H. pylori and A2 and B subunit genes of Vibrio cholerae. Also claimed are chimeric proteins (see AAW80599-600) encoded by such recombinant DNAs, methods for the recombinant production of the chimeric proteins, and use of the chimeric proteins in preventative and therapeutic vaccines for H. pylori associated diseases such as gastritis, gastric ulcer, duodenal
                                                                                                                                                                                                                                                                                                                                                                      New chimeric proteins for use against Helicobacter pylori - comprising antigenic protein of H. pylori and Al and B subunits of Vibrio cholerae toxin, preferably produced by recombinant techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAR-1997;
31-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine; antigen; antigen; toxin; diagnosis; gastritis; ulcer; stomach cancer; babB gene; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
Helicobacter pylori.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Helicobacter pylori babB gene PCR primer
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                                                                                                                                                                Sequence 22 BP; 9 A; 4 C; 4
                                                                                                                                                                                            ulcer and gastric cancer
                                                                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-568279/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kim B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9844130-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DAEW-) DAEWOONG PHARM CO LTD.
                                                                                                                      Local
                                                                                 2015 CAGGGGATGGGAAAAAACCTT 2035
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                                                                                                                                                                                                                                                                                                                                               2-21; Page 15; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shin
                                                                                                                         Similarity
                                                       CCGTGGATGAAAAAAAACCTT 21
                                                                                                              Conservative
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97KR-00011951
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                        0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                           Score 14.6; D
Pred. No. 2.4e
0; Mismatches
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                                                                                                                                                                  5 T; 0 U; 0 Other;
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                                                                                                                          .4e+03
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                                                                                                                                         Length
                                                                                                               Indels
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AAT99486
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                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a purified peptide having at least one of CC the following properties: (a) it stimulates transcription of a DNA CC sequence under the control of a delta A element, an NPKB element or an CC Ikaros binding oligonucleotide consensus sequence; (b) it binds to any of a delta A element, an NPKB element or an Ikaros binding oligonucleotide consensus sequence; (c) it competitively inhibits the binding of a calement or an Ikaros isoform to any of a delta A element, an NPKB element, an NPKB element, an NPKB competitively inhibits the binding of a competitively inhibits the binding of a competitively inhibits the control competitively inhibits protein protein interactions of transcriptional complexes formed with naturally occurring Ikaros isoforms. The proteins, provided that they stimulate gene transcription under the control of delta A elements, NPKB elements and/or Ikaros-binding oligonucleotides, bind to delta A elements, NPKB elements and/or Ikaros-binding oligonucleotides, binding of a helments, NPKB elements and/or Ikaros-binding oligonucleotides, binding to inhibit vely inhibit Ikaros binding to Ikaros-responsive elements and/or inhibit protein interactions of transcriptional complexes with the control of the competitively inhibit Ikaros binding to Ikaros-responsive elements and/or inhibit protein interactions of transcriptional complexes with
                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 17
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14-SEP-1993;
02-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                naturally occurring Ikaros isoforms, can be used to treat immune system disorders, e.g. leukaemia or AIDS, or corpus striatum disorders, e.g. Alzheimer's disease. AAV66975 to AAV67118 represent oligonucleotides
                                                                                                                                                                                                                                                                                                                                                   Sequence
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Mus
ST receptor; heat stable toxin receptor; colorectal cancer; tumour;
                                Human ST receptor PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Col 26; 111pp; English.
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Synthetic.
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                                                                   21-MAY-1998
                                                                                                                                 AAT99486 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or corpus striatum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ikaros poly:peptide(s) - useful for treating disorders of immune system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-582621/49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                          5690 TACCACTGTTTTGCCTTCCTT 5710
                                                                                                                                                                                                                                                                                                                                                                                      H
                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                   1 Similarity
                                                                                                                                                                                                                                                                                                                                                                                  the present
                                                                                                                                                                                                                                                                                                                                                   22 BP;
                                                                                                                                                                                                                     TTCCCCTGTTTTGGTTTCCTT
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                                                                                                                                                                                                                                                                                   Conservative
                                                                 (first entry)
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93US-00121438.
94US-00238212.
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                                                                                                                                                                                                                                                                                                                                                     12
                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                   A; 3 C; 7
                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88
                                                                                                                                                                                                                                                                                                                                                                                    invention
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                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                   Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                   G; 0
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system; corpus striatum; AIDS;
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                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                  1; Length 22;
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                                                                                                                                                                                                                                                                                   Gaps
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Claimed PCR primers (see AAT99462-T99531) hybridise to sequences that cencode the extracellular domain of human heat-stable toxin (ST) receptor protein (see AAW37371), a highly specific marker for metastasised colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in AAT99486 and AAT99487. Claimed in vitro methods for determining whether or not (i) an individual has metastasised colorectal cancer cells, or (ii) a tumour cell is a colorectal cancer cells, or (iii) a tumour cell is a colorectal cancer cells for tumour cell sensitive straintestinal tissue and/or body fluids or tumour cells from an individual to determine whether ST receptor protein is being expressed by cells in the sample. Expression is determined by immunoassay or by PCR using primers that selectively amplify ST receptor cDNA or mRNA (see also cancer colls and the sample of colorectal cancer cells amplify ST receptor cDNA or mRNA (see also cancer colls and colorectal cancer cells amplify ST receptor cDNA or mRNA (see also cancer cells and colorectal cancer cells amplify ST receptor cDNA or mRNA (see also cancer cells and colorectal cancer cells amplify ST receptor cDNA or mRNA (see
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Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                            toxin receptor on cells in a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                Determining whether an individual has cells and origin of tumour cells - by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-008454/01
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22
BP; 7 A; 1 C; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Carrithers SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JEFFERSON THOMAS
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G; 4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                metastasised colorectal
detecting presence of he
0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     of heat-stable
                                                                                                                                                                                                                                                                                                   receptor
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Ś Query Match Best Local Matches 3199 АСТСАССССТТСАСАЛАСТС 3219 0.2%; 1 Similarity 81.0%; 17; Conservative AATGAGGGCTGGAAATAGTG 0; Score 14.6; DB 1; Pred. No. 2.4e+03; Mismatches DB 1; Length Indels 22; 0; Gaps

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RESULT 3370
AAT99484
ID AAT9948
XX AAT9948
XC AAT9948
XX ID HUMAN S
XX FI FECE
KW ST FECE
KW METASTA
DE HOMO 98
XX W SYNThet
OS HOMO 98
XX W W097425
XX W097425
                 02-MAY-1997;
                                                       13-NOV-1997.
                                                                                                                                                Synthetic
                                                                                                                                                                                   ST receptor;
metastasis; o
                                                                                                                                                                                                                                       Human ST receptor PCR primer.
                                                                                                                                                                                                                                                                                                                    AAT99484;
                                                                                                                                                                                                                                                                                                                                                      AAT99484 standard; DNA; 22 BP
                                                                                          WO9742506-A1
                                                                                                                                                                                                                                                                               21-MAY-1998
                                                                                                                             sapiens
                                                                                                                                                                                   ; heat stable toxin recordiagnosis; human; PCR;
                                                                                                                                                                                                                                                                             (first entry)
                   97WO-US007467
                                                                                                                                                                                                    receptor; colorectal cancer; tumour;
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AAT9949
ID AAT9949
XX AAT9949
XX AAT9949
XX 21-MAY-
XX TI rece
XW ST rece
XW ST rece
XW METASTA
COS HOMO B8
XX SYNTHE
COS HOMO B8
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Best Local
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Determining whether an individual has metastasised colorectal cancer cells and origin of tumour cells - by detecting presence of heat-stal toxin receptor on cells in a sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ST receptor; heat stable toxin metastasis; diagnosis; human; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1998
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                                                                                                      1998-008454/01.
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; heat stable toxin receptor; colorectal diagnosis; human; PCR; primer; ss.
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                                                                                                                                                       SL,
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; tumour;
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encode the extracellular domain of human heat-stable toxin (ST) receptor protein (see AAW37371), a highly specific marker for metastasised colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in AAT99494 and AAT99495. Claimed in vitro methods for determining whether or not (i) an individual has metastasised colorectal cancer cells, or (ii) a tumour cell is a colorectal cancer cell comprise the steps of examining a sample of extraintestinal tissue and/or body fluids or tumour cells from an individual to determine whether ST receptor protein is being expressed by cells in the sample. Expression is determined by immunoassay or by PCR using primers that selectively amplify ST receptor cDNA or mRNA (see also AAT97229)
                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 14; Page 53; 62pp;
                                                                                                                                                                                                                                                                                                                                                                                     (вее
                                                                                                                                                                                                                                                                                                                                                                                     AAT99462-T99531) hybridise to sequences that
                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
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Matches Query Match Best Local : Sequence Similarity 22 BP; 7 Conservative A; 1 C; 10 G; 4 T; 0 U; 0 Other; 0.2%; 0 Score 14.6 Pred. No. Mismatches 14.6; .4e+03 DB 1; Length Indels 0

Gaps

0

RESULT 3372 AAT99490 AAT99490; AAT99490 standard; DNA; 22 ВP

밁 5

3201 TGAGGGGCTTGAGAAAGTGGG 3221

2 TGAGGGGCTGGAAATAGTGAG 22

Homo ST receptor; heat stable toxin receptor; colorectal cancer; metastasis; diagnosis; human; PCR; primer; ss. Synthetic 21-MAY-1998 sapiens. TS receptor (first entry) PCR primer. tumour;

03-MAY-1996; 02-MAY-1997; 13-NOV-1997. 96US-0016564P 97WO-US007467

(UYJE-) UNIV JEFFERSON THOMAS Carrithers SL;

WPI; 1998-008454/01. Waldman SA,

Determining whether an individual has cells and origin of tumour cells - by toxin receptor on cells in a sample. metastasised colorectal cancer detecting presence of heat-stat of heat-stable

14; Page 53; 62pp; English.

Claimed PCR primers (see AAT99462-T99531) hybridise to sequences that encode the extracellular domain of human heat-stable toxin (ST) receptor protein (see AAM37371), a highly specific marker for metastasised colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in AAT99490 and AAT99491. Claimed in vitro methods for determining whether or not (i) an individual has colorectal cancer receptor

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RESULT 3373
AAT99496
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Best Local S
Matches 17
Query Match
Best Local Similarity
                                                                                 Claimed PCR primers (see AAT99462-T99531) hybridise to sequences that encode the extracellular domain of human heat-scable toxin (ST) receptor protein (see AAW37371), a highly specific marker for metastasised colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in AAT99497 and AAT99497. Claimed in vitro methods for determining whether or not (i) an individual has metastasised colorectal cancer cells, or (ii) a tumour cell is a colorectal cancer cells, or (iii) a tumour cell is a colorectal cancer cells of examining a sample of extraintestinal tissue and/or body fluids or tumour cells from an individual to determine whether ST receptor protein is being expressed by cells in the sample. Expression is determined by immunoassay or by PCR using primers that selectively amplify ST receptor cDNA or mRNA (see also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ST receptor; heat stable toxin receptor; colorectal cancer; tumour; metastasis; diagnosis; human; PCR; primer; ss.
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                                             Sequence
                                                                                                                                                                                                                                                                                             Claim 14; Page 53; 62pp; English.
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AAT97229)
                                                                         usıng primers
AAT97229)
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 0.2%;
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                                             1 C;
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 Score 14.6;
Pred. No. 2.
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Pred. No. 2
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                DB 1;
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                                             Other;
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                                                                                                                                                                                                                                                                                                                                        presence of heat-stable
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                22;
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2.4e+03;

RESULT 3375

AAT99492

AAT99492

standard; DNA;

22

ВP

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RESULT 3374
AAT99488
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                                                                                                                                                                                                                           colorectal cancer cell comprise the steps of examining a sample of extraintestinal tissue and/or body fluids or tumour cells from an individual to determine whether ST receptor protein is being expressed by cells in the sample. Expression is determined by immunoassay or by PCR using primers that selectively amplify ST receptor cDNA or mRNA (see also
                                                                                                                                                                                                                                                                                                                                     colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in ANT99488 and ANT99489. Claimed in vitro methods for determining whether or not (i) an individual has metastasised colorectal cancer cells, or (ii) a tumour cell is a
                                                                                                                                                                                                                                                                                                                                                                                                                                             Claimed PCR primers (see AAT99462-T99531) hybridise to sequences that encode the extracellular domain of human heat-stable toxin (ST) receptorein (see AAW37371), a highly specific marker for metastasised
                                                                                                                                                                   Sequence 22
                                                                                                                                                                                                           using primers
AAT97229)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Determining whether an individual has metastasised colorectal cancer cells and origin of tumour cells - by detecting presence of heat-stal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ST receptor; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Waldman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-NOV-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9742506-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human ST receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT99488 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                     Local
                                     3199 AGTGAGGGCTTGAGAAAGTG 3219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3201 TGAGGGGCTTGAGAAAGTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor on cells in a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17;
N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGAGGGGTTGGAAATAGTGAG
                                                                                                                                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis; human; PCR; primer; ss.
                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Carrithers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       heat stable toxin receptor; colorectal cancer; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-0016564P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-US007467
                                                                                                                                                                   7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer.
                                                                                                   0.2%;
                                                                                                                                                                 1 C; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                       Score 14.6;
                                                                                                     Pred.
                                                                                                                                                                 G; 4 T; 0 U; 0 Other;
                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                     No.
                                                                                                     4e+03
                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.
                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                           22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of heat-stable
                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   receptor
                                                                                 Gaps
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RESULT 3376
AAV27972
ID AAV2797
XX
AC AAV2797
XC AAV2797
XX
DT 25-SEP-
XX
DE Ataxia
XX
KW 88; PCF
KW radiati
XX
CS Synthet
OS Homo 88
                                                                                                                                                                                                                                                                                                                                                                       밁
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                           protein (see AAW37371), a highly specific marker for metastasised colorectal cancer cells. PCR using these primers provides specific and sensitive detection of human ST receptor expression. A specific primer pair comprises the primers given in AAT99492 and AAT99493. Claimed in vitro methods for determining whether or not (i) an individual has metastasised colorectal cancer cells, or (ii) a tumour cell is a colorectal cancer cell comprise the steps of examining a sample of extraintestinal tissue and/or body fluids or tumour cells from an individual to determine whether ST receptor protein is being expressed by cells in the sample. Expression is determined by immunoassay or by PCR using primers that selectively amplify ST receptor cDNA or mRNA (see also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claimed PCR primers (see AAT99462-T99531) hybridise to sequences that encode the extracellular domain of human heat-stable toxin (ST) recep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Determining whether an individual has cells and origin of tumour cells - by toxin receptor on cells in a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-NOV-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ST receptor; heat stable toxin receptor; colorectal cancer; tumour; metastasis; diagnosis; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human ST receptor PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 14; Page 53; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-008454/01.
                                                                                        Ataxia telangiectasia exon 8 primer 1.
                                                                                                                                                                           AAV27972 standard;
                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Waldman SA,
   Homo sapiens
                                               radiation;
                                                                                                                      25-SEP-1998
                                                                                                                                                   AAV27972;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYJE-) UNIV
                                                           PCR; primer; amplification; ataxia telangiectasia; diagnosis; human;
                                                                                                                                                                                                                                                                                 3199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens.
                                                                                                                                                                                                                                                                                                             1 Similarity
17; Conserv
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                                                                                                                                                                                                                                                                        AGTGAGGGGCTTGAGAAAGTG 3219
                                                                                                                                                                                                                                                      AATGAGGGGCTGGAAATAGTG 21
                                                                                                                                                                                                                                                                                                                                                                     22 BP; 8
                                               breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carrithers SL;
                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JEFFERSON THOMAS
                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-0016564P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-US007467
                                                                                                                                                                                                                                                                                                                                                                     A; 1 C; 9
                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                22
                                                                                                                                                                                ₽P
                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                     G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                           Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  metastasised colorectal cancer detecting presence of heat-stable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detecting presence
                                                                                                                                                                                                                                                                                                                            4e+03;
                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                          Length 22
                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptor
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RESULT 3377
AAV16099/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The primers AAV27964-V28086 are used to amplify ataxia telangiectasia (ATM) exons and their adjacent splice junction sites. These can be used as a method of detecting a mutation in the ATM gene by comparing the PCR products of amplification from a sample from a patient suspected of having an ATM mutation with a sample from a non-mutated ATM patient This method is especially useful for diagnosing ataxia telangiectasia in heterozygotes and can be used to locate the positions of the mutation. The diagnosis of ataxia telangiectasia in patients needing therapeutic radiation will prevent fatal radiation burns and the development of breast cancer which can occur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 4 A; 1 C; 6 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method of detecting ataxia telangiectasia - comprises use of primers based on intron-exon boundaries, useful for diagnosing disease in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-NOV-1997;
                                                                                                                                                                                                                 Glutathione-S-transferase-UL97 fusion Protein; GST; amplification; PCR; CMV; protein kinase.
                                                                                                                                                                                                                                                                              07-JUL-1998
                                                                                                                                                                                                                                                                                                   AAV16099;
                                                                                                                                                                                                                                                                                                                           AAV16099 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 6; Page 5; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     heterozygotes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-312503/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9822621-A1
                                                                                                        25-JUL-1997;
                                                                                                                                05-FEB-1998.
                                                                                                                                                                                                                                                      pGST97-delta-N238 primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (VIRG-) VIRGINIA MASON
                                                          (HARD)
                                                                                26-JUL-1996;
                                                                                                                                                        WO9804747-A1
                                                                                                                                                                               Human herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                      6469 TTTTTCTGTTTGTGAATAAGG 6489
                                                                                                                                                                                                                                                                                                                                                                                                                                     il Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                        1 TTTTTCTGTATGGGATTATGG 21
                                                          HARVARD
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                              (first entry)
                                                                                  96US-0022888P
                                                                                                        97WO-US013340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-00753147.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97WO-US020953.
                                                          COLLEGE
                                                                                                                                                                                  ຫ
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RES CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   .4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                88;
                                                                                                                                                                                                                               primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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WPI; 1998-130727/12.

Coen D;

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RESULT 3378
AAX81911
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The pGST97-delta-N238 primer was used with pGST97 reverse primer (CC (AAV16101) to amplify the corresponding fragments by PCR, using CC (Inearised pGST-UL97 as a template. The PCR product would encode for a CC glutathione-S-transferase (GST)- human cytomegalovirus (CMV) UL97 fusion CC protein in which the N-terminal region of the latter protein was CC truncated by 238 residues. Truncation of the N-terminal region of UL97 (CC protein kinase (see also AAV16100) and Carried out to CC investigate any changes in enzyme activity. It was determined that CC truncation of the first 303 N-terminal residues totally abolished CC enzymatic activity indicating that the N-terminal region was involved in CC kinase activity. The invention provides an assay method for the detection CC fagents that inhibit or enhance activity of the CMV UL97 protein CC kinase. GST-UL97 fusion protein confers an advantage in this method as it is more soluble, without loss of enzymatic activity, than UL97 protein. CC the invention claims that by using an optionally modified polypeptide which contains a UL97 phosphorylation consensus sequence, the assay contains that by using an optionally modified polypeptide which contains a UL97 phosphorylation consensus sequence, the assay contains that the variance of the compositions which can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                    Vaccine; T cell receptor; TCR; T cell; V beta 6.2/3; V beta 6, V beta 6.7; V beta 2; V beta 5/1; V beta 7; V beta 13; V beta multiple sclerosis; PCR primer; ss.
                                                                                                                                                                                                                                 03-DEC-1997;
                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX81911 standard; DNA;
The specification describes vaccines which comprise immunologically
                                                                                                                          Brostoff
                                                                                                                                                         (XIMM-)
                                                                                                                                                                                                    03-DEC-1997;
                                                                                                                                                                                                                                                               10-JUN-1999.
                                                                                                                                                                                                                                                                                             WO9927957-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-SEP-1999
                               Example 10;
                                                                                            WPI; 1999-404801/34.
                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Assay for detecting modulating agents of UL97 protein kinase - useful for, e.g. preparation of therapeutics for treatment of cytomegalovirus
                                                                                                                                                                                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 Similarity
17; Conserv
                                                                                                                                                       IMMUNE
                                                            receptor
                                                                                                                          WS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CGGCCCGGGCCCGGGATCCGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGAGCCCGGCCCGGGATCCCG 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              treating CMV infections
                            Page 41; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                       Wilson
                                                                                                                                                       RESPONSE CORP.
CANCER CENT SIDNEY.
                                                                                                                                                                                                    97WO-US023147.
                                                                                                                                                                                                                                 97WO-US023147
                                                                                                                                                                                                                                                                                                                                                                                                                                  to amplify human
                                                            peptide-derived vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P
                                                                                                                       DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                       Smith LR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.6;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 1 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                TCR V beta genes.
                                                                                                                          Gold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .4e+03;
                                                                                                                       ₽P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.
                                                                                                                          Carlo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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Matches
                                                         Query Match
                                                                                                                                                                                           than passively administering heterologous antibodies, allows the host's own immune system to mobilize and suppress auto aggressive T cells. Therefore, the suppression is persistent and may involve any and all immunological mechanisms in effecting that suppression. Such a multifaceted response is more effective than the uni-dimensional suppression achieved by passive administration of monoclonal antibodies or extantderived regulatory T cell clones. PCR primers AAX81882-X81914 were used to amplify and analyse human TCR V beta genes, in the course of the
                                                                                                                                                                                                                                                                                                                                                                                                                       effective amounts of T cell receptor (TCR) peptides. The TCRs are present on the surface of T cells. The TCRs are chosen from V beta 6.2/3, V beta 6.6/5, V beta 6.7, V beta 2, V beta 5/1, V beta 7 or V beta 13. The V beta TCRs peptide-based vaccines are useful for prevention or treatment of multiple sclerosis. The presence of V beta 6.7 appears to be particularly associated with multiple sclerosis and can be used to determine an individual's susceptibility to multiple sclerosis. Vaccinating, rather
                                                                                                                   Sequence
                                  Local
l Similarity
17; Conserv
                                                                                                                22
     Conservative
                                                                                                                BP; 4 A;
                            0.2%;
                                                                                                                6
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     0; Mismatches
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Pred. No. 2
                                                                                                                7 T; 0
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Human leukocyte antigen; HLA; allele; HLA-B*3913; HLA-B*1406; human; HLA-B*51; HLA-DRB1*0820; HLA-DRB1*04; HLA-DRB4*01; allele typing; exon; major histocompatibility complex; MHC; PCR primer; ss.
                                                                                                                                                                            WO9954496-A2
                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                    HLA allele DRB1*0820 exon 2 amplifying primer.
                                                                                                                                                                                                                                                  11-FEB-2000
                                                                                                                                                                                                                                                                           AAZ39210 standard; DNA;
                                                                                                                                                                                                                                                                                                                  4141 CTGTGTACCTGATTTGTTCTC 4161
                                                                                                                                                                                                                                                                                                           μ
                                                                                                                                                                                                                                                                                                           CAGTGACCCTGAGTTGTTCTC
                                                                                                                                                                                                                                                  (first entry)
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The invention provides polynucleotides corresponding to exon 2 and exon 3 of human leukocyte antigen (HLA) alleles HLA-B*3913, HLA-B*1406 and HLA-B*51 and exon 2 of HLA alleles HLA-DBB1*0820, HLA-DRB1*04 and HLA-DRB4*01. The polynucleotides are useful for typing the above HLA alleles in a sample, especially by a method that comprises (a) amplifying all/part of the relevant sequence using at least one primer pair; and (b) hybridizing the amplified product to a set of probes specifically hybridizing to target regions comprising one or more polymorphic nucleotides of the sequence, to determine the absence or presence of the

New polynucleotides for human leukocyte antigen, useful for typing HIA alleles

HLA,

allele

fragments,

for typing HLA alleles

WPI; 1999-634008/54.

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20-APR-1998; 19-APR-1999; 28-OCT-1999.

98EP-00870088 99WO-EP002614.

(INNO-) INNOGENETICS

Claim 4;

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English.

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RESULT 3380
AXX88451/c
ID AAX8845
XX AXX8845
XX DT 01-OCT-
XX PANTES;
XW RANTES;
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XW 29-JUL-
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XW RANT
This invention describes novel oligonucleotides which are used for detecting the chemokines RANTES (regulated upon activation normal T expressed and secreted), macrophage inflammatory protein (MIP)-1 alpha or MIP-1 beta by (a) obtaining a sample possible containing RANTES or MIP-1 calpha or MIP-1 beta RNA, (b) performing an isothermal transcriptional camplification on the sample with 2 oligonucleotide primers, (c) detecting the product of step (b) where detection of a product indicates the presence of RANTES, MIP-1 alpha or MIP-1 beta in the sample. The assay is completed to determine the levels of the chemokines RANTES, MIP-1 alpha and the sample of the chemokines range. The assay is to be inhibitors of CD4+ T-cells by primary non-syncytium-inducing HIV-1 strains. Thus the level of expression of these genes can be used as prognostic markers for direct therapeutic management of HIV-infected patients. By being isothermic, the assay requires less manipulation by the experimenter. Also 'spiking' the sample with a known amount of control RNA allows quantitation and qualification of the products in a single assay. AAX88447-X88491 represent the primers and probes used inthe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RANTES; chemokine; detection; primer; probe; amplification; MIP-1 alpha; regulated upon activation normal T expressed and secreted; MIP-1 beta; macrophage inflammatory protein; CD4+ T-cell; inhibitor; prognosis; primary non-syncytium-inducing HIV-1 strain; therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detection of expression levels of the cytokines RANTES, MIP-lalpha MIP-lbeta used as prognostic markers of HIV-infected patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human RANTES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 38; 48pp;
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                                                                                                                                                                                                         establishment of paternity. Kits containing the products of the invention provide many copies of selected RNA targets under conditions of constant temperature, ionic strength and pH. Specific amplification of RNA targets
                                                                                                                                                                                                                                                             This invention describes a novel method for the autocatalytic amplification of an RNA target in a transcription-based amplification system without thermalcycling. The method generates oligonucleotides for diagnostic or environmental testing, for use e.g. as probes and in cloning. Typical applications are the detection of genetic or infectious diseases, the monitoring of responses to therapy, the quantitation or detection of microorganisms in foods, forensic studies and the establishment of paternity. Kits containing the products of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Autocatalytic amplification; transcription-based amplification; thermalcycling; diagnostic; environmental testing; probe; detect genetic disease; infectious disease; microorganism; food; forens
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                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kit for autocatalytic amplification of RNA targets
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                                                                                                                                                                                     increases sensitivity, convenience,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               690 CCTGGATGTGGCCATGAGGCA 710
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17; Conserv
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90US-00550837.
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                                                                                                           G; 3 T; 0 U;
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                                                                                                                                                                                           reliability of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
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RESULT 3382

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RESULT 3383
AAX34398
ID AAX3439
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AC AAX3439
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                                                                                                                                                                                                                                                            The present sequence represents a PCR primer for human APRIL (a CC proliferating inducing agent). APRIL is a member of the tumour necrosis CC factor family, and essentially free of normally associated proteins. CC APRIL and APRIL antibodies are useful in pharmaceutical compositions for preventing or reducing severity of an autoimmune disease or an immune CC response to tissue graft. The composition is also useful for stimulating the immune system, and treating cancer. APRIL is also CC useful for treating APRIL-related disorders by delivering via a vector (preferably viral vector) (gene therapy) into a mammalian (preferably CC human) cell. Labeled APRIL and fragments are useful for intifying APRIL CC receptors by screening compositions. Antisense DNA and antibodies and CC modified APRIL (preferably an anti-APRIL receptor antibody) are useful as blocking agents for inducing cell death by interfering with APRIL CC, and treats, suppresses or alters an immune response involving a signalling pathway between APRIL and its receptor (preferably involving a CC signalling pathway between APRIL and its receptors (preferably involving a CC also administered, and radiation therapy is also given to the patient is also administered, and radiation therapy is also given to the patient
                                                                                                                                                                        Query Match
Best Local S
Matches 17
   AAX34398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A Proliferating Inducing Agent (APRIL), a member Factor Family - useful as diagnostic agents and ftreatment of immune disorders and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-1997;
26-MAR-1998;
                               AAX34398 standard; DNA; 22 BP
                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 30; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-215028/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human APRIL PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX30126 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APRIL; tumour necrosis factor; TNF; proliferating inducing agent; immune disorder; cancer; PCR primer; ss.
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                                                                                                                                  5651 CCAGCCTCATCCTCTTAGTTG 5671
                                                                                                                                                                         17;
                                                                                                            -
                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                       22 BP; 2
                                                                                                                                                                         Conservative
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98US-0079384P.
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                                                                                                                                                                                       Score 14.6; DB 1; Length 22; Pred. No. 2.4e+03;
                                                                                                            21
                                                                                                                                                                                                                                     8 T; 0 U; 0 Other;
                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                       AAX32149
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Palmer L, F.
                                                                                                                                                                                                                                                                                                                                                                                                This sequence represents a probe used to detect the coding region for a novel 3-hydroxyacyl-CoA dehydrogenase (hcd) from Staphylococcus aureus (AAX34393). The products can be used to prevent or treat bacterial infection, e.g. S. aureus infection or Helicobacter pylori infection. They can also be used for preventing invasion of bacteria in damaged tissue including wounds in skin or connective tissue caused, e.g. by mechanical, chemical, thermal or radiation damage or by implantation of indwelling devices, or wounds in the mucous membranes, such as the mouth the mucous membranes, such as the mouth the mucous membranes.
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                                             Allele profile; diagnosis; treatment; pharmacogenetic; breast cance: CFTR; cystic fibrosis; dystrophin; Duchenne muscular dystrophy; p53 Becker muscular dystrophy; hi-Fraumeni syndrome; neurofibromatosis; colorectal cancer; MSH2 gene; MLH1 gene; BRCA1 gene; BRCA2 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue; wound; skin; connective tissue; implant; mutous membrance; throat; mammary gland; urethra; vagina; probe; hybridisation; ss.
          Synthetic
                                    colorectal
BAP1 gene;
                                                                                                             BRCA2 gene
                                                                                                                                        14-JUN-1999
                                                                                                                                                                  AAX32149;
                                                                                                                                                                                           AAX32149 standard;
                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated 3-hydroxyacyl-CoA dehydrogenase polynucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-263995/22.
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Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3-hydroxyacyl-CoA dehydrogenase; hcd; infection; Helicobacter pylori; tissue; wound; skin; connective tissue; implant; mucous membrane; mou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S.aureus 3-hydroxyacyl-CoA dehydrogenase gene
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                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                              2158 ATCCAATTCTACAAGTCCACC 2178
                                   gene;
                                                                                                                                                                                                                                                                                                           l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Page 55; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BRIGHAM & WOMENS HOSPITAL VIRUS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                       mammary glands, urethra or vagina
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                                    cancer;
PCR prim
                                                                                                              specific primer.
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                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                        (first
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                                                                                                                                                                                           DNA;
                                                                                                                                       entry)
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                                    2 gene; MLH1 gene; BRCA1 gene; BRCA2 ss.
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                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                     Score 14.6;
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                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                         2.4e+03
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                                                                                                                                                                                                                                                                                                                                                                0 Other;
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                                                                                                                                                                                                                                                                                                                                    Length
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                                                                                                                                                                                                                                                                                                             Indels
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                                                                                     breast cancer,
                                                                                                                                                                                                                                                                                                                                      22;
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                                                                                                                                                                                                                                                                                                                                                                                                    the mouth,
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RESULT 3385
AAZ09309/c
ID AAZ0930
XX Human m
XX Macroph
KW Macroph
KW Macroph
KW Chemoth
XX Haematc
KW Chemoth
XX Haematc
XX Homoo se
XX US59488
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XX US59488
XX T16-DEC-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to methods of determining a functional allele CC profile of a gene in a population. Functional allele profiles comprise the commonly occurring alleles in a population, and the relative CC trequencies at which such alleles of a given gene occur. The methods are CC used to identify and determine the frequency of the functional alleles of Genes which display extensive allelic heterogeneity, particularly those CC implicated in disease or conditions, such as the BRCA1 gene associated CC with breast cancer, CFTR associated with cystic fibrosis, dystrophin CC associated with Duchenne muscular dystrophy and Becker muscular GC dystrophy, and p53 associated with Li-fraumeni syndrome. The methods can CC also be employed for diseases where allelic and genetic heterogeneity CC exist, such as breast cancer, neurofibromatosis, and hereditary non-CC polyposis colorectal cancer. Identification of functional alleles is necessary for identification of mutations which may be implicated in the CC disease. Sequences AAX32001-172 represent primers for determining the functional allele profiles of various genes. The primers are specific for genes such as MSH2 gene, MLH1 gene, BRCA1 gene, BRCA2 gene and BAP1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-AUG-1997;
22-MAY-1998;
                       16-DEC-1996;
                                                         07-SEP-1999.
                                                                                                                                                      Synthetic
                                                                                                                                                                                     Macrophage stimulating protein, MSP; human; modulator; proliferation; differentiation; intestinal epithelium; colon crypt; treatment; cance haematopoietic disorder; megakaryovyte deficiency; gastrointestinal; chemotherapeutic agent; gut toxicity; PCR primer; ss.
                                                                                                                                                                                                                                                                                Human macrophage stimulating protein PCR primer 1.
                                                                                                                                                                                                                                                                                                                                                                                              AAZ09309 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                 sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTTTGAGATTTGTGTTTGTGT 7327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCTTAGATTTGTGTTTTGGT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 13 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               common functional alleles in a population - use f disease associated with allelic heterogeneity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                      (first entry)
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98US-00084471.
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                       96US-00766982
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Pred. No. 2.4e+03;
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Best Local S
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                                                                                                                                                                                               product of the invention binds to RON (a cell membrane protein tyrosine kinase which is a member of the c-met family) to promote the formation of colon crypts. MSP analogues are useful for the treatment of conditions requiring the administration of MSP, such conditions include haematopoietic disorders such as those involving a deficiency of megakaryocytes and gastrointestinal disorders such as ulcerative colitis, Crohn's disease and infections. The MSP analogues are useful for cancer, where the aggressive use of chemotherapeutic agents or the use of cancer, where the aggressive use of chemotherapeutic agents or the use of whole body radiation may lead to gut toxicity. The MSP analogues, which have a higher activity than normal human MSP are effective at smaller dosages, or optionally, they may be administered less frequently than human MSP. This sequence represents a PCR primer used to amplify the human MSP described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel purified and isolated analogue of mature macrophage stimulating protein (MSP) having at least one unpaired cysteine residue substituted with another amino acid which modulates the proliferation or differentiation of the intestinal epithelium. The
                                                                                                                                                           Sequence 22 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or haematopoietic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analogues of macrophage stimulating protein for treating gastrointestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-517975/43
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                             2826 TTCCAAGCCCCAGGAGCTGTG 2846
21 Trechedacechedecerrere 1
                                                                            1 Similarity
17; Conserv
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                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96US-00766982.
                                                                                                                                                             A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                 0.2%;
                                                                            0
                                                                                                 Score 14.6;
Pred. No. 2
                                                                              Mismatches
                                                                                                   .4e+03
                                                                                                                    DB 1;
                                                                                                                    Length
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RESULT 3386
AAZ27795/c
ID AAZ2779
XX
AC AAZ2779
XX
AC AAZ2779
XX
Tandem
KW Tandem
KW Tandem
KW Tandem
KW DNA typ
XW DNA t ITR sequence; pentanucleotide tandem repeat; stutter artifact; DNA typing; DNA profiling; linkage analysis; criminal justice; paternity testing; animal lineage analysis; microsatellite locations. AAZ27795 standard; DNA; Schumm JW, AAZ27795; 04-FEB-1998; 04-FEB-1999; 12-AUG-1999 Synthetic. Tandem repeat sequence; DNA isolation; intermediate tandem repeat; PCR primer 23-DEC-1999 WO9940194-A1 Homo sapiens polymorphism (PROM-) PROMEGA CORP. for human DNA marker clone G212. Bacher JW; detection; (first 98US-00018584 99WO-US002345 entry) 22 PCR primer; 88 ΒP

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PPI XX PFX PFX XX PFX XX PFI XX PFI XX PFX XX PFI X
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AAX21479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC fragments, at least one of which contains an ITR sequence, a region of CC the DNA fragment which contains at least one repeat unit consisting of a CC sequence of five, six or seven bases repeated in tandem at least two CC times; (b) providing a stationary support having at least one CC oligonucleotide associated with it, where the oligonucleotide includes a CC sequence of nucleotides which is complementary to a portion of the ITR CC sequence; and (c) combining the DNA fragments with the support under CC conditions where the DNA fragments including the DNA fragment containing CC used to isolate DNA containing pentanucleotide tandem repeat sequences as CC well as to detect target ITR DNA sequences having a low incidence of CC stutter artifacts (no more than 2.4%). The method is useful in DNA CC profiling for linkage analysis, criminal justice, paternity testing and CC other forensic and medical uses. DNA typing is also useful for confirming CC the lineage of horses, dogs and other prize animals. The invention CC overcomes problems related to the use of microsatellite loci in DNA CC profiling. The method can detect polymorphisms with a low incidence of the lineage of forest, which has previously been a repolation intermenting confirming.
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Best Local
Martinez Herrera
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                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX21479;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               stutter artifacts, which has previously been a problem in interpret allelic content of loci. The development of markers based on larger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence method of the
                              Ponz Ascaso F,
                                                                                                                                                                                                                                           21-JAN-1999.
                                                                                                                                                                                                                                                                                               WO9902718-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           repeat units, enables easier separation of the fragments on electrophoretic gels. This allows the simultaneous analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolating DNA DNA profiling.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-590696/50
                                                                                                                                    09-JUL-1997;
                                                                                                                                                                                      09-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hybridization
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2861
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
                                                                                                                                                                                                                                                                                                                                                                                                                            virus;
                                                                                                                                                                                                                                                                                                                                                 mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                   INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer for Turnip mosaic virus genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGAAAGCAAGCAGGTGGCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGGAAGCAAGGAGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA containing intermediate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                        amplification;
                                                                                NACIONAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                    virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intermediate tandem repeat (ITR) sequence using selection, and comprises: (a) providing several
                                                                                                                                                                                                                                                                                                                                                                                                                               epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention. The method is for isolating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               is a PCR primer for a
                                                                                                                                  97ES-00001522
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                           Torres
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; 9 C; 3
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                                                                             INVESTIGACION & TECNOLOGIA
                                                                                                                                                                                                                                                                                                                                                                                                                            tion; infection; plant; Arabidopsis thaliana; edible vaccine; ss.
                        Pascual V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22
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Pred. No. 2.
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                              Sanchez Sanchez
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tandem repeat sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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AAV83869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-120919/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rumour suppressor gene; IB3089A; DBCCR1; 9q32-33; hypermethylation; Deleted in Bladder Cancer Chromosome Region candidate 1; cancer; bladder cancer; ovarian cancer; skin cancer; squamous carcinoma; renal carcinoma; squamous cell ocspice; squamous carcinoma; renal carcinoma; squamous cell ocspice; squamous carcinoma; renal carcinoma; squamous cell ocspice; squamous carcinoma; squamous cell ocspice; squamous carcinoma; squamous cell ocspice; squamous carcinoma; squamous cell ocspice; squamous cell ocspice; squamous carcinoma; squamous cell ocspice; squamous cell ocspice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  method to construct the complete turnip mosaic virus (TuWV) genome. The invention relates to the isolation of an infectious clone of TuWV. The clones, and vectors containing the plants, specifically Arabidopsis thaliana, for basic virological research and for expression of genes or epitopes of interest, particularly expression of animal virus epitopes for production of edible vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22
                                                                                                                                                                                                                New human gene for diagnosis,
                                                                                                                                                                                                                                                                                                                              Knowles M,
                                                                                                                                                                                                                                                                                                                                                                                                                   28-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prenatal testing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reverse primer SSCPex4 used to amplify IB3089A gene coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV83869;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   296 GCATTGGCACTGTGGGGAAAC 316
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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                                                                                                                                                                                                                                       gene,
                                                                                                                                                                                                                                                                                                                              Habuchi
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                                                                                                                                                                                                                                                                                                                                                                                                                      97GB-00010995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-GB001515
                                                                                                                                                                                                                                       DBCCR1, associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    heterozygosity loss; PCR primer;
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No. 2
                                                                                                                                                                                                                   bladder and of cancer.
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PCR primers AAV83864-83 were used to amplify the coding region from the exons of the genomic DNA of tumour suppressor gene IB3089A, also known a DBCCR1 (Deleted in Bladder Cancer Chromosome Region candidate 1). The gene is located at 1932-33. Although the gene is expressed in multiple numan tissues, lack of expression of the gene was found in several bladder cancer cells. The DBCCR1 gene is involved in the development of

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Disclosure;

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PR 12-NON
PR 12-NON
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PR 22-MAN
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PR WPI;
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07-NOV-1997;
12-NOV-1997;
01-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and/or hypermethylation of the remaining allele, leading to reduced expression of the gene. DBCCR1 nucleic acid sequences are used for treatment or prevention of cancer, particularly bladder, ovarian or skin cancer, squamous carcinoma, renal cell carcinoma or squamous cell oseophageal carcinoma. The DBCCR1 protein can be used directly in the products can be used for diagnosis or prognosis of cancer, or to indicate prediaposition to cancer (including prenatal testing), particularly where this is associated with loss of heterozygosity at 932-33. Antibodies may also be used for affinity purification, therapeutically as modulator and to detect the protein in cells etc
The present invention describes genomic DNA which contains a BRCA2 gene where the first 12 nucleotides beginning exon 5 are 5'-TCCTGTTGTTC-3' a in sequence (I) (see AAX03249), where nucleotides numbers 5782-5790 are GTTTGTGTT as in sequence (IV) (see AAX30255), and where the last 20 nucleotides encoding exon 15 are 5'-CTGCGTGTTCTCATAAACAG-3' as in sequence (II) (see AAX30251) and the first 20 nucleotides beginning exor 16 are 5'-CTGTATACGCGTTTC-3' as in sequence (III) (see AAX30253). Products and methods from the present invention can be used for identifying mutations in the BRCA2 gene leading to predisposition or
                                                                                                                                                                                                                   New coding sequence haplotypes of the human BRCA2 gene - used to develop products for determining susceptibility to, detection and treatment of breast or ovarian cancer.
                                                                                                                                                                                                                                                                                                                                                Murphy PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9909164-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human BRCA2 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX33028 standard;
                                                                                                                                                                                                                                                                                            WPI; 1999-190163/16.
                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                   (ONCO-) ONCORMED INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
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                                                                                                                                                                                                                                                                                                                             'n,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B₽;
                                                                                                                                                                                                                                                                                                                                                White MB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genetic testin breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-0055784P.
97US-0064926P.
97US-0065367P.
98US-00071715.
                                                                                                                                                                                                                                                                                                                           Schryer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer SEQ ID NO:41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                226pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         testing; protein therapy; haplotype; detection;
cancer; ovarian cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                               'n
                                                                                                                                                                                                                                                                                                                                                Rabin MB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
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                                                                                                                                                                                    English
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                Olson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 Other;
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                                                                                                                                                                                                                                                                                                                                                  Yoshikawa
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                                                                                        Query Match
Best Local (
                                                                           Matches
                                                                                                                                                                                                   higher susceptibility to breast or ovarian cancer. They can also be used for detection and gene therapy for breast and ovarian cancers. They can be used in methods for monitoring disease progression, for determining patients suited for gene and protein replacement progression, or for detecting the presence or quantifying the amount of a tumour growth inhibitor following such therapy. The BRCA2 protein, polypeptides, their functional equivalents, antibodies, and PNs may also be useful in the study of the characteristics of BRCA2 proteins, such as structure and function of BRCA2 in oncogenesis or subcellular localisation of BRCA2 protein in normal and cancerous cells. AAX33001 to AAX33097 represent PCR
                                                                                                                                                  Sequence 22 BP; 13 A; 5 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                       primers used in the amplification of the human BRCA2 gene
                                     7307
22
                                                                           17;
                                                                                            Similarity
 CTCTTAGATTTGTGTTTTGGT 2
                     CTTTGAGATTTGTGTTTGTGT 7327
                                                                           Conservative
                                                                                            0.2%;
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Pred.
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RESULT 3390
AAX36878
                                             Human XLIS
                                                                                          AAX36878;
                                                                                                                AAX36878 standard;
SCLH; cortical
                       XLIS gene;
                                                                   14-JUL-1999
          issencephaly;
                     human; detection; diagnosis;
                                             gene fragment PCR primer 2.1
                                                                    (first
                                                                                                                 DNA;
                                                                    entry)
                                                                                                                  22
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genetic Synthetic. neurodegenerative sapiens. counselling; an; detection; diagnosis; prenatal diagnosis; therapy; LIS; agyria-pachygyria; subcortical laminar heterotopia; dysgenesis; cryptogenic epilepsy; neurological disorder; the disease; Alzheimer's disease; X-linked disorder; primer; ss

EP918091-A1

26-MAY-1999 97EP-00402811

21-NOV-1997; 97EP-00402811.

(INRM) INSERM INST NAT SANTE & RECH MEDICALE

WPI; 1999-290318/25.

Chelly J,

Kahn

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Portes

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Pinard J;

New gene and its gene product expressed in the brain, useful diagnosing and treating disorders such as lissencephaly and diagnosing laminar her heterotopia. subcortical for

9; Page 46; 71pp; English

the XLIS gene transcripts or to compare their sequence with that of the XLIS transcripts from patients for in vitro especially prenatal diagnosis of lissencephaly (LIS) (or agyria-pachygyria), subcortical laminar heterotopia (SCLH), cortical dysgenesis, cryptogenic epilepsies or neurodegenerative diseases such as Alzheimer's disease. These disorders mainly affect females as the XLIS gene is X-linked. The XLIS fragments may also be used to administer to patients to prevent or treat the above disorders and may be used as a tool in genetic counselling. This sequence is a primer for the human XLIS gene of the invention. The XLIS fragments may be used to detect abnormalities in the expression of the XLIS gene transcripts or to compare their sequence with that of the

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AAX81831/c
 Query Match
Best Local S
Matches 17
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Best Local
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AAX81831
                                                                               PCR primers AAX81830-31 were used to amplify human malignancy-associated gene (MAG) proteins. The polypeptide is useful for detecting antibodies associated with liver disease. Probes derived from the MAG gene are useful for detecting the presence of sequences associated with neoplastic disease, e.g. liver diseases such as cirrhosis and hepatocellular carcinoma, and therefore can be used in disease diagnosis. The sequences can be used for development of therapeutics that are useful for inhibition of the development of neoplastic liver disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 XIIS gene from a sample for comparison to normal samples in the in vitro diagnosis regime. This may also be performed by amplifying XIIS cDNA from the mRNA in the sample. Antibodies to XIIS may be used to detect XIIS in a biological sample or can be administered to patients to prevent or treat the above disorders. They may also be used to purify XIIS from a biological sample, XIIS may also be administered to patients to prevent or treat the above neurological disorders. In addition XIIS may be used or treat the above neurological disorders. In addition XIIS may be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX81831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         as a marker of neuronal cells at an early stage of development; its discovery increases understanding of both the neuronal movement which leads to development of the cortical region of the brain and of the pathogenesis of the group of neuronal disorders mentioned above
                                                                                                                                                                                                                                               Liver-associated malignancy-associated gene for cirrhosis and hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                               12-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Liver neoplastic disease; malignancy-associated neoplastic disease; cirrhosis; hepatocellular ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                          Sequence
                                                                                                                                                                                                                                                                                           WPI; 1999-404942/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer used to amplify human malignancy-associated gene (MAG).
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 l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
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                                                                                                                                                                                                                                                                                                                    Ljubimova JY,
                                                          BP;
                                                                                                                                                                                                                   Page 17;
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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             0.2%;
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                                                        7 C; 4
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                                                                                                                                                                                                                                                                                                                      Demetriou AA;
                                                                                                                                                                                                                   English.
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                                                        G; 7 T;
Score 14.6; D
Pred. No. 2.4e
0; Mismatches
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                       0 U;
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U;
             2.4e+03
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                             DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 Other;
                                                                                                                                                                                                                                                           (MAG), useful for screening
                             1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene; MAG;
                          Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                           neoplastic
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RESULT 3392
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KW neoplag
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XX Synthet
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PT for ci
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AAA99208
ID AAAS
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AC AAAS
AC AAAS
AC AAAS
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DJ 23-C
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DE Huma
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                                                                                                                                                                                                                                                                                                             RESULT 3393
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The specification describes a liver neoplastic disease polynucleotide and malignancy-associated gene (MAG) proteins. The polypeptide is useful for detecting antibodies associated with liver disease, Probes derived from the MAG gene are useful for detecting the presence of sequences associated with neoplastic disease, e.g. liver diseases such as cirrhosis
                                                                                                                                                                                                         AAA99208;
                                                                                                                                                                                                                                                            AAA99208 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           associated with neoplastic disease, e.g. liver diseases such as cirrhosis and hepatocellular carcinoma, and therefore can be used in disease diagnosis. The sequences can be used for development of therapeutics that are useful for inhibition of the development of neoplastic liver disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Liver-associated malignancy-associated gene (MAG), for cirrhosis and hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Black K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Liver neoplastic disease; malignancy-associated gene; MAG; liver disease; neoplastic disease; cirrhosis; hepatocellular carcinoma; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer used to amplify cDNA sequences isolated from liver tissue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-SEP-1999
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                                                                                                                                                      23-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primers AAX81836-37 were used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                         616
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTGTGAGCTGGCGAATGCTG 636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 23; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No. 2
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Human; apoptosis; CCR9; anti-tumour;

tumour;

cancer; diagnosis; primer;

Human apoptosis related protein CCR9 related primer #4.

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RESULT 3394
AAZ39687/C
ID AAZ3968
XX AAZ3968
XX AAZ3968
XX CAAZ3968
XX CAAZ3968
XX Gene po
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes the human CRR9 protein, which is an apoptosis related protein having apoptosis inducing activity. Human CCR1 has anti-tumour activity, and can to produce a malignant tumour gene treating agent. The CCR9 gene and protein can be used for the diagnosis of the resistance of cancer cells against an anticancer agent. The present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example from the present sequence represents a primer which is used in an example of the present sequence present sequence represents a primer which is used in an example of the present sequence present seque
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gene polymorphism; human; Vth aggregation factor; genetic diagnosis; diabetes; FPCR; SSCP; fluorescence-based polymerase chain reaction; single strand conformation polymorphism; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Vth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ39687 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22 BP; 10 A; 6 C; 4 G;
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                                                       Disclosure; Page 10; 34pp; Japanese
                                                                                                          Discrimination of human V aggregation
                                                                                                                                                            WPI; 2000-057352/05
                                                                                                                                                                                                                                                                                                                                                                                                                      JP11313676-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                                                                                                               30-APR-1998;
                                                                                                                                                                                                                                                                                                                 30-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                    16-NOV-1999
                                                                                                                                                                                                          (SAKA ) OTSUKA PHARM CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present sequence represents a primer which present invention
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   factor gene specific FPCR-SSCP primer.
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Pred. No. 2.
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                                                                                                          factor
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                                                                                                       gene polymorphism
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The invention provides a method for the discrimination of the gene

conformational polymorphism) analysis

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AAZ49922
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymorphism of human Vth aggregation factor, where one of the following (1) to (6)) residues/nucleotides in the aggregation gene is discriminated in the patient to be tested: (1) residue 495: guanine (G) or adenine (A), (2) residue 642: (G) or thymine (T), (3) residue 263: (G) or (A), (4) residue 2763: (G) or (A), (5) residue 2863: (A) or (G), (6) residue 5112: (A) or (G). The method is useful in the genetic diagnosis of a diabetes patient. The method uses FPCR-SSCP (fluorescence-based polymerase chain reaction-single strand conformation polymorphism) for analyzing DNA samples for polymorphisms. Sequences AAZ39632-717 represent primers used for the FPCR-SSCP analysis of the human Vth aggregation factor gene
        referred as DBCCRI (Deleted in Bladder Cancer Chromosome Region candidate 1), and its promoter in a tumour suppressor region at 9932-33 between D9S1848 and AFMA239XA9 of human chromosome 99. The DBCCRI sequence can be used in the diagnostic, prophylactic and therapeutic treatment of cancer particularly bladder cancer. The promoter can be used to design DBCCRI amplifying primers and screen compounds that activate production of DBCCRI It can also be used in methods for determining inactivation of the DBCCRI gene in a patient, where inactivation indicates the presence of a tumour or a predisposition to cancer, especially a cancer associated with loss of heterozygosity involving 9932-33 e.g. bladder cancer, squamous carcinoma, skin cancer, renal cell carcinoma, oesophageal and ovarian cancers. The present sequence is a reverse primer SSCPex4 used to amplify exon 4 of IB3089A from the genomic DNA for SSCP (single-stranded conformational no light of the property of the present sequence is a reverse primer SSCPex4 used to amplify exon 4 of IB3089A from the genomic DNA for SSCP (single-stranded conformational no lights) and proverse primer sychematics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; tumour suppressor; IB3089A; cytostatic; promoter; 9q32-33; beleted in Bladder Cancer Chromosome Region candidate 1; DBCCR1; diagnostic; prophylactic; therapeutic treatment; cancer; skin; over bladder; squamous carcinoma; renal cell; oesophageal; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnostic; prophy bladder; squamous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human tumour suppressor gene IB3089A exon 4 reverse primer SSCPex4
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                                                                                                                                                                                                                                                                                The patent relates to the identification of a novel gene IB3089A, also referred as DBCCR1 (Deleted in Bladder Cancer Chromosome Region candidate
                                                                                                                                                                                                                                                                                                                                                 Claim 22; Page 78; 101pp;
                                                                                                                                                                                                                                                                                                                                                                                            cancer or
                                                                                                                                                                                                                                                                                                                                                                                                              Novel promoter sequence, useful for identifying a predisposition
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-171014/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Knowles M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (IMCR ) IMPERIAL CANCER RES TECHNOLOGY
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                                                                                                                                                                                                                                                                                                                                                                                       the presence of a tumor.
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AAZ44367
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                                RESULT 3397
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Best Local S
Matches 17
                                                                                       Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                     This invention describes a novel human G-protein coupled receptor protein capable of binding lysophosphatidic acid, and proteins derived from it by addition, deletion and/or substitution of one or more amino acid residues. Antibodies to the protein are used for diagnosis of, and agonists/antagonists to the protein are used for the treatment of, prostatic disorders such as prostate cancer, benign prostatic hyperplasia, and inflammation of the prostate. This sequence represents a primer used in the isolation of the human G protein-coupled receptor protein described in the method of the invention
AAA11724;
                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                        G protein-coupled receptor; treatment; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                AAA11724 standard; DNA; 22
                                                                                                                                                                                                                       Example 1; Page 60; 67pp; Japanese.
                                                                                                                                                                                                                                              G-protein coupled receptor protein binding to lysophosphatidic acid used
                                                                                                                                                                                                                                                                                                                                                                WO9967383-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                Human G protein-coupled receptor primer IO6R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ44367 standard; DNA; 22
                                                                                                                                                                                                                                                                                Nozaki Y,
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                                                                                               Score 14.6;
Pred. No. 2.
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Pred. No. 2.4e+03;
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                                                                                       Mismatches
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AAA95380
                                                                                                                                                                                                                                                                    RESULT 3398
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                    AAA95380;
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  Parkinson's
                         Rat; Nurr1;
                                                                      Rat Gli1 coding sequence PCR primer #1.
                                                                                                                      12-FEB-2001
                                                                                                                                                                                                                    AAA95380 standard;
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tyrosine hydroxylase; catecholamine-related disease; disease; manic depression; schizophrenia; PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequences. The Nurr1 protein is involved in the induction of tyrosine hydroxylase expression in adult rat-derived hippocampal progenitor cells. The Nurr1 gene and protein can be used in the treatment of catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia. They can also be used to induce tyrosine hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such a
                          Inhibiting cell proliferation useful for cancer therapy, comprises administering Rad51 inhibitor in vivo.
                                                                                                                                                                                                                                                           10-FEB-1999;
06-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             radiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense inhibitor; human; Rad51; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human Rad51
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                                                                                                          WPI; 2000-506091/45.
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99US-00454495.
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Pred. No. 2
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AAA74138
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents an antisense inhibitor of human Rad51, designated AS6 (also referred to as R51AS6). The antisense inhibitors can be used in a method of the invention, for inhibiting cell proliferation. They can also be used in methods for inducing sensitivity to radiation and DNA damaging chemotherapeutics in an individual and in a method for prolonging survival in an individual with cancer. The methods and antisense molecules are useful for inhibiting cell proliferation, especially cancerous cell proliferation, for inducing sensitivity to radiation and DNA damaging chemotherapeutics in individuals and for prolonging survival in an individual with cancer. Kits for carrying out addition that the cancer and for the second contents that the cancer and for the second carrying out and the cancer that the cancer and for the second carrying out addition that the cancer and for the cancer and for the cancer and for the cancer and for the cancer and th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer; loblolly pine; Simple Sequence Repeat; SSR; microsatellite DNA repeat; genetic marker; mapping; inheritance population genetics study; plant breeding programme; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22 BP; 6 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                             polynucleotide having simple sequence repeat useful as markers in plants for genetic characterization e.g. genetic mapping study, an inheritance study of a commercially important trait in a plant breeding program.
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                                                                                                                                                                                                                                                                                                                            WPI; 2000-482836/42.
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19-JAN-1999;
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                                                                                                                                                                                 Claim 6; Page 24; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                             (ECHT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                        ) INT PAPER CO.
) ECHT C S.
) NELSON C D.
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                                                                                                                                                                                                                                                                                                                                                                                                                    NELSON C D.
US SEC OF AGRIC.
                                                                                                                                                                                                                           tic characterization e.g. genetic a commercially important trait in
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nilarity 81.0%;
Conservative
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99US-00232785.
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No. 2
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The present invention relates to loblolly pine polynucleotides with one or more Simple Sequence Repeats (SSRs) (see AAA74205-A74322). SSRs are also known as microsatellite Day Repeats. The SSRs are useful as genetic markers for genetic mapping, population genetics studies and inheritance studies in various plant breeding programmes. The present sequence is a PCR primer used for detecting the presence of a SSR locus in a pine

genomic DNA sample

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AAS01199
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Matches
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                                                                                                                Query Match
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                                                                                                                                                            The sequence represents the human Rad51 antisense oligonucleotide, AS6. The antisense oligonucleotide is used to study down-regulation of Rad51 protein in human brain, breast and prostate cells. Rad51 protein is defective in repair of damaged DNA, genetic recombination and the recombinational repair of DNA lesions, and plays a central role in cancer. Inhibiting cell proliferation involves administering to a cell a Rad51 antibody or its fragment. The Rad51 antibody or its fragment is useful for inhibiting cell proliferation, for treating disease states such as cancer, autoimmune disease, arthritis, graft rejection, inflammatory bowel disease, proliferation induced after medical procedures such as surgery, angioplasty etc. in humans and animals
                                                                                                                                                                                                                                                                                                                                        Inhibiting cell proliferation for treating arthritis, graft rejection, inflammatory bowel disease, cancer, proliferation induced after medical procedure, involves administering Rad51 antibody or its fragment to cel
                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-244704/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-SEP-1999;
06-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; Rad51; antisense; drug screening; cancer; autoimmune disease; arthritis; graft rejection; inflammatory bowel disease; surgery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human RAD51 antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS01199 standard; cDNA; 22 BP
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                                                                                                                                                                                                                                                                                                                   Example 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                               CCCTAGTAATACTTAAGGCAC 3497
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                                                                                                                                                                                                                                                                                                                 Fig 16C; 102pp; English.
                                                                                                                                        BP; 6
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99US-00455300.
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                                                                                                                                        9 C; 3
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                                                                                                                                        G; 4 T; 0 U; 0 Other;
                                                                                                   Score 14.6; DB 1;
Pred. No. 2.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.6; DB 1;
Pred. No. 2.4e+03;
0; Mismatches 4
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                                                                                                                Length 22;
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                                                                                        Indels
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RESULT 3403
AAC92055/c
ID AAC9205
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AC AAC9205
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AC AAC9205
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AC AAC9205
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RESULT 3402

AAC92055; 21-MAR-2001

(first entry)

AAC92055 standard;

DNA;

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                                                                   Query Match
Best Local
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                                                                                                                               The invention relates to antibody-based fusion proteins having one or more mutations in the junction between an immunoglobulin (Ig) and a no Ig moiety, which increases the circulating half-life of the fusion protein. The serum half-life of the mutant fusion protein is improved without affecting the interaction of the antibody moiety with the Fc receptor (FcR) and Fc protection receptor (FcRp). The antibody-based fusion proteins of the invention are useful to treat cancer, viral infections, immune disorders, and to enhance growth (including proliferation) of specific cell types. The present sequence is an oligonucleotide used for generating mutant humanised KS antibody (recognises epithelial cell adhesion molecule) and interleukin-2 fusion protein with Pro to Leu substitution at the fusion junction
                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                           Example
                                                                                                                                                                                                                                                                                                                                                   Antibody-based fusion protein comprises mutations near the fusion junction for enhancing circulating half-life.
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-514646/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-FEB-2001; 2001WO-US004455
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fusion protein; immunoglobulin; serum half-life; Fc protection receptor; FcRp; cancer; viral infectel proliferation; human; KS antibody; EpCAM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Epithelial cell adhesion molecule; mutagenesis; interleukin-2; IL-2; ds
                        6998 GGGAAAGGGAGATTTTCTTCT 7018
                                                      1 Similarity
17; Conservat
N
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                                                                                                           BP; 5 A;
                                                      Conservative
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/note= "The 5' end of the complementary strand
the 3' end of this sequence by 5'-CGC-3'"
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/label=
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                                                                  0.2%;
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                                                                                                           4 C; 9 G; 4 T; 0 U;
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                                                                  Score 14.6; DB 1;
Pred. No. 2.4e+03;
22
                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       infection;
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                                                                             Length
                                                      Indels
                                                    0;
                                                                                                                                                                                                                                                                                 and a non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disorder;
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                                                    Gaps
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RESULT 3404
AAF58870/c
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Best Local
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New nucleic acid encoding
                                                                                                                                                                                                                                                      AAF58870 standard;
                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for determining the ability of a test compound to alter the circadian rhythm of a mammal by its ability to alter phosphorylation or degradation of one or more period proteins. The present sequence is a PCR primer for one such period protein (rat PER1), that was used in the method of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Keesler G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rat; PER1; circadian rhythm;
                   WPI; 2001-258133/26.
                                                                                                                                                                                            Human metastasis-associated antigen C4-4A PCR primer
                                                                                                                                                                                                                  06-JUN-2001
                                                                                                                                                                                                                                     AAF58870;
                                                                                               29-SEP-2000; 2000WO-EP009567
                                                                                                                                     WO200123553-A2
                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10;
                                                         (DEKR-) DEUT
                                                                            29-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying compounds that alter circadian rhythm of a mammal by altering the phosphorylation or degradation of human period proteins, comprises adding test compound to screening system comprising period proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AVET )
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                                                                                                                                                                                                                                                                                                                          3239
                                                                                                                                                                           human;
                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                            1 Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                       TTGTCAGCAGCCTTAACCAGA 1
                                                                                                                                                                                                                                                                                                                                                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 33; S5pp; English.
                                                                                                                                                                            metastasis-associated antigen; C4.4A; cancer;
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                                                                                                                                                                                                                                                                                                                                             Conservative
                                      Roesel M,
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                                                          KREBSFORSCHUNGSZENTRUM
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                                                                             99US-00407784
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                                                                                                                                                                                                                                                       DNA;
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                                      Wuerfel
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                                                                                                                                                                                                                                                        ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              period protein; PCR primer;
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                              Mismatches
human metastasis-associated antigen
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                                                                                                                                                                                                                                                                                                                                                              DB 1;
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RESULT 3405
AAF58876/c
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Best Local S
Matches 17
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Best Local 9
                                                                                                                 The present invention provides the protein and coding sequences of the human and rat metastasis-associated antigen C4.4A. The protein is expressed rarely in the adult, except on metastasising cancer cells. Because of this, the sequences are useful in cancer diagnosis and treatment of cell proliferation diseases. The present sequence is a PCR primer used to isolate the human C4.4A coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein and coding sequences of the human and rat metastasis-associated antigen C4.4A. The protein is expressed rarely in the adult, except on metastasising cancer cells. Because of this, the sequences are useful in cancer diagnosis and treatment of cell proliferation diseases. The present sequence is a PCR primer used to isolate the human C4.4A coding sequence
                                                                                                                                                                                                                        New nucleic acid encoding rat or human metastasis-associated antigen {\sf C4.4A} for treating cell proliferative disorder associated with a metastasizing tumor.
                                                                                                                                                                                                                                                                                                                                                                                                 05-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 BP; 6 A; 12 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          metastasizing tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C4.4A for treating cell proliferative disorder associated with
                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                 29-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                       WO200123553-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat; human; metastasis-associated antigen; C4.4A; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human metastasis-associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF58876;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF58876
                                                                                                                                                                                                   Example 1; Page 31; 63pp; English
                                                                                                                                                                                                                                                                                                   Zoeller M,
                                                                                                                                                                                                                                                                                                                                                                        29-SEP-2000; 2000WO-EP009567
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
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                                            l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
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criredadcerederedere
                      CTTTGGGGAATGGGGTGGGGG
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                                                                                                                                                                                                                                                                                                  Roesel M,
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                                              Conservative
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                                                                                            12 C; 2 G;
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                                                                                                                                                                                                                                                                                                   Wuerfel J;
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                                                          Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.6;
Pred. No. 2
                      3631
                                                                                             2 T; 0 U;
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                                              Mismatches
                                                          14.6;
No. 2
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                                                                     DB 1;
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AAF99703
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AC AAF9970
AAF99
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AAF58878/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the protein and coding sequences of the human and rat metastasis-associated antigen C4.4A. The protein is expressed rarely in the adult, except on metastasising cancer cells. Because of this, the sequences are useful in cancer diagnosis and treatment of cell proliferation diseases. The present sequence is a primer used to isolate the human C4.4A coding sequence
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                                                                          Synthetic
                                                                                                                                             Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                                                                              AAF99703 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid encoding rat or human metastasis-associated antigen C4.4A for treating cell proliferative disorder associated with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-SEP-2000; 2000WO-EP009567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rat; human; metastasis-associated antigen; C4.4A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human metastasis-associated antigen C4-4A PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-JUN-2001
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                            WO200122972-A2
                                                                                                                                                                                                                                                   Immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                    12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                     AAF99703;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   metastasizing tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-258133/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200123553-A2
                                                                                                                           infectious
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTTTGGGGAATGGGGTGGGGG 3631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTTTGGAGCGTGGGTGGGTG
                                                                                                                        disease; allergy; immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roesel M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-00407784.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 C;
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Pred. No. 2.
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                                                                                                                      deficiency; phosphorothioate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    #6
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RESULT 3408
AAH79377/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence aphosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response. The method comprises administering an immunostimulatory nacid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vaccinating against tumors, infe using immunostimulatory Py-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (IOWA )
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27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2000; 2000WO-US026383
           мао у,
                                                              29-OCT-1999;
                                                                                       29-OCT-1999;
                                                                                                                  11-JUL-2001.
                                                                                                                                            CN1302893-A.
                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                               Human; RNA uncoiling enzyme 44; cancer; nervous system disease; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                      Human RNA uncoiling enzyme 44 coding sequence
                                                                                                                                                                                                                                                                  04-DEC-2001
                                                                                                                                                                                                                                                                                                                     AAH79377 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Krieg AM,
                                     (BODA-)
                                                                                                                                                                                                                                                                                            AAH79377;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                   5328
                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                                                                                                                                                        μ.
                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
17; Conserv
                                     BODAO
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           Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                  CTCTCTTTGCCTCACTCTCTC 5348
                                                                                                                                                                                                                                                                                                                                                                                        CTCTCTCTCTCTCTCTCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schetter C, .
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                  (first entry)
                                     GENE TECHNOLOGY CO
                                                                99CN-00119928.
                                                                                         99CN-00119928
                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         81.0%;
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                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        infectious diseases, allergies rich and TG nucleic acids.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.6;
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                                     SHANGHAI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                         PCR primer #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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WPI; 2001-566052/64.

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RESULT 3409
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Best Local :
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                         PCR primers AAH74499-AAH74500 were used to amplify a fragment of the pig genome, comprising the marker SW782. The primers were used in the method of the invention. The specification describes a method for determining if a pig is predisposed to a boar taint. The method comprises assaying for the presence of alleles conveying susceptibility to boar taint using genetic markers selected from SW1057, SW782, S0121, SW222 or regions of chromosome 6 spanning in between, or SW857, SW2496, SW295, SW210, S0007, SW761, SW1557 or regions of chromosome 14 spanning in between. The method is useful for determining the predisposition of pigs to boar taint which is a strong unpleasant odour given off upon heating or cooking of meat from uncastrated male pigs, and for exhibiting desirable flavour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polypeptide-human RNA uncoiling enzyme 44 and polynucleotide for coding it.
                                                                                                                                                                                                                                                              Determining if a pig is predisposed to boar taint for exhibiting desirable flavor properties, involves assaying for the presence alleles conveying susceptibility to boar taint using specific ge
                                                                                                                                                                                                                                                                                                                                                         Haley CS,
                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000; 2000GB-00002451
                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-FEB-2001; 2001WO-GB000448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chromosome 14; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pig; boar taint;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer used to amplify a marker SW782 from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH74500
                                                                                                                                                                                                                                                                                                                             WPI; 2001-496928/54
                                                                                                                                                                                                                    Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention provides the protein and coding sequences of human uncoiling enzyme 44. The sequences can be used in the treatment of cer and nervous system diseases. The present sequence is a PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            coding sequence of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTTTCAAAAAACAAATTTTCC 1
                                                                                                                                                                                                                                                                                                                                                         Archibald
                                                                                                                                                                                                                 Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 14 (Disclosure);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nt; genetic marker;
SW857; SW2406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                   32; 71pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SW210;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pig genome
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AAH78021

AAH78021 standard; DNA;

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RESULT 3410
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Best Local
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Best Local S
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                                                 Matches
                                                                                                                                                        PCR primers AAH74330-41 were used to amplify a fragment of the human ATIP gene. ATIP has isoforms designated hATIP2, hATIP3, hATIP4, hATIP5 and hATIP6. All ATIP proteins comprise in their C-terminals a common fragment which interacts with the angiotensin II (AT2) receptor. ATIP proteins have antionogenic functions. The human ATIP gene has 17 exons, and is have antionogenic functions. The human ATIP gene has 17 exons, and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 6 A; 1 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                         New protein family, designated hATIP, which interacts with the AT2 receptor of angiostatin II are anti-oncogenic and useful to detect treat cancer or precancerous conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ATIP; hATIP2; hATIP3; hATIP4; hATIP5; hATIP6; AT2 receptor;
angiotensin II receptor; antioncogenic; 8p21.3-p22; cancer; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                     07-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                           07-FEB-2001; 2001WO-FR000359
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                                                                                               Sequence
                                                                                                                        condition,
                                                                                                                                  located at chromosome region 8p21.3-p22. ATIP polypeptides are used to detect, evaluate or g
                                                                                                                                                                                                                                    Claim 7; Page 100; 118pp; French.
                                                                                                                                                                                                                                                                                                                                  Nahmias C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200157209-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer
                                                                                                                                                                                                                                                                                                                                                            (CNRS ) CNRS
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                        7401 AGCAAGCAACATCAGCAGCAG
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N
                                                 l Similarity
17; Conserv
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AACAAGACACATAAGCAGCAG
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                                                                                                                         and as an
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                                               0.2%;
llarity 81.0%;
Conservative
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                                                                                               11 A; 5 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                        anti-tumour medicament
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                                                             Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fragment of the human ATIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.6;
Pred. No. 2
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                                                 Mismatches
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                                                             4e+03
                                                                       DB 1; Length 22;
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                                                                                                                                  P polynucleotides
give prognosis fo
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                                                  Indels
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                                                  <u>,</u>
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                                                 Gaps
                                                                                                                                      cancer
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AAC84905
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primers AAH78021-22 were used to amplify cDNA encoding a human alpha subunit of prolyl 4-hydroxylase, designated alpha (III) subunit. The alpha (III) subunit is useful for the production of recombinant collagen, and for the diagnosis, prevention and treatment of various diseases and disorders associated with decreased or increased production of the subunit in specific tissues. The polynucleotide is a source of probes and primers, which are useful for diagnosis, prevention and treatment of diseases and disorders associated with increased or decreased expression or activity of various prolyl 4-hydroxylase enzymes and to identify alpha or beta prolyl 4-hydroxylase or fragments in tissue, e.g. biopsies from specific tissues, etc or other biological samples. Small molecules that modulate, regulate and inhibit prolyl 4-hydroxylase activity are useful for treating and preventing kidney fibrosis and various other diseases
                                                    SECX; cytostatic; gynecological; gene therapy; screening assay; human; chromosomal mapping; forensic biology; cell proliferation; cancer; cell differentiation; immune associated disorder; gestational disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New alpha subunit of prolyl 4-hydroxylase and polynucleotide encoding the subunit, useful for diagnosis, prevention and treatment of diseases and disorders associated with increased or decreased expression of the
                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 52; 75pp;
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collagen; kidney fibrosis; PCR primer; ss.
             Homo sapiens
                                                                                                               Primer Ag
                                                                                                                                                                                                     AAC84905 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         subunit.
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                                          PCR primer;
                                                                                                                                                                                                                                                                                                                                   1 Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                81.0%;
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Pred. No. 2.
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                                                                                                               expression analysis
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                                                                                                                                                                                                                                                                                                                                                                                             8 T; 0 U; 0
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                                                                                                                                                                                                                                                                                                                                                                                             Other;
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                                                                                                               of SEC1
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AAS11868/c
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoding them. The SECX polypeptides can be expressed by standard recombinant methodology. The SECX polypeptides are useful for treating or preventing a SECX-associated disorder. The invention is useful in screening assays; detection assays (e.g. chromosomal mapping, cell and tissue typing, forensic biology); predictive medicine (diagnostic assays, prognostic assays, monitoring clinical trials, and pharmocogenomics); and methods of treatment (e.g. therapeutic and prophylactic), especially disorders characterized by aberrant cell proliferation and/ or differentiation like cancer or immune associated disorders or gestational disease. Sequences AAC84904-906 represent primer-probe sets employed in the expression analysis of SEC1 clone
                                                                                                                       Homo sapiens
Synthetic.
                                                                                                                                                          Human; PCNA; PCR primer; HIS tag; iminodiacetic acid cellulose; IDAC; immobilised protein; proliferating cell nuclear antigen; ss.
                                                                                                                                                                                                Human PCNA random mutagenesis PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polynucleotides encoding SECX proteins useful for treating characterized by an aberrant level of cell proliferation ar differentiation like cancer or immune associated disorders.
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22-JUN-2000;
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20-JUL-1999;
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                        12-MAR-1999;
                                              12-MAR-1999;
                                                                                                US6232083-B1
                                                                                                                                                                                                                          24-OCT-2001
                                                                                                                                                                                                                                                 AAS11868;
                                                                                                                                                                                                                                                                        AAS11868 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to human SECX polypeptides and polynucleotides
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                                                                       15-MAY-2001
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17; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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2000US-00604286
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                       9908-00268536
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99US-0144722P.
                                                9905-00268536
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                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 9 C; 2 G; 6 T; 0 U;
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                                                                                                                                                                                                                                                                          ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.6;
                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length
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VIND (ANAM)

NEW YORK STATE RES

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RESULT 3414
AAF76175/C
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in genetically engineered cells, transferring the protein to metal charged iminodiacetic acid cellulose (IDAC), immobilising the protein on the metal charged IDAC and detecting the protein or its functional activity immobilised on the IDAC. The method is useful for determining the expression of a protein and functional activity of a protein, which includes binding specificity, enzyme activity, stimulation of enzyme activity or stimulation of delta-polymerase activity. The protein is especially human or Drosophila melanogaster proliferating cell nuclear antigen (PCNA). Metal charged IDAC allows easy screening of a large number of proteins following mutagenesis and can rapidly ascertain which mutants have desired functional activity or binding capacity. The present
                                                                                                                                                                                                                                                                           Transgenic mouse; immunodeficient; tissue recipient; lymphocyte deficient; human cytokine; interleukin; II-7; II-6; SCF; LIF; stem cell factor; leukaemia inhibitory factor; GM-CSF; M-CSF; granulocyte macrophage-colony stimulating factor; macrophage-colony stimulating factor; mapor histocompatibility complex; allergenicity determination; human monoclonal antibody generation; haematopoietic cell development; human immune system animal model; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to determining expression or functional activity of a protein, comprises expressing the protein having a polyhistidine region in genetically engineered cells, transferring the protein to metal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Determining protein expression and function by genetically expressing proteins having polyhistidine regions, immobilizing the protein on metal charged iminodiacetic acid cellulose and detecting protein or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF76175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF76175 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Col 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activity.
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               WPI; 2001-169001/17
                                              Huang MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                Human M-CSF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence is a mutagenic PCR primer for random mutagenesis of
                                                                                                               31-AUG-1999;
                                                                                                                                                                                                                WO200115521-A1
                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                               30-AUG-2000; 2000WO-US023971.
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                                                                                GENENCOR INT INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGGAAGAAGGGTCTAAATC 4763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGATGAAGAAGGATCTTAAGC 2
                                              Harding FA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer, SEQ ID NO:41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zaika
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                               99US-0151688F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 C; 2 G; 8 T; 0 U; 0 Other;
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human PCNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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New transgenic mice, useful as non-human mammalian models of human disease, comprise recombination activation gene mutations and donor specific transgenes encoding cytokines.

CC both alleles of a gene activated in early lymphocyte development are conditional activated in early lymphocyte development are conditional activated in early lymphocyte development are conditional activation gene-2 (RAG-2) gene are conditional activational gene-2 (RAG-2) gene activational generate the conditional generate generate the conditional generate generate fully human polyclonal conditional antibodies to generate fully human polyclonal conditional generate generate generate the generate the generate the generate the generate the conditional generate generate generate the generate ge immunologically dysfunctional. In addition the transgenic mouse provides a unique model system which supports T cell development in a manner which more closely resembles normal ontogeny, as they possess CD4+ T cells in the periphery that exhibit MHC-restricted antigen- specific responses. Sequences AAF76133-AAF76192 represent human cytokine PCR primers used in the development of human cytokine-expressing transgenic mice The invention relates to a transgenic immunodeficient recipient mouse which is capable of supporting the growth of donor cells. In the mous the mouse,

Sequence 22 BP; 7 A; 6 C; 6 G; 3 T; 0 U; 0 Other; DB 1;

Matches Query Match Best Local (Similarity Conservative 0.2%; 0, Score 14.6; Pred. No. 2 Mismatches .4e+03; Length 22; Indels 0 Gaps 0

4803 CTGCCCTTGTATGACCCGGAT 4823

CIGCCCTIGIAAGACTIGGCT 2

밁 S

22

RESULT 3415 AAF31837/c

AAF31837 standard; DNA; 22 BP

12-APR-2001 (first entry)

Human MAT II beta subunit-specific PCR primer.

Human; methionine adenosyltransferase II; MAT II; MAT II beta subunit; cytostatic; immunosuppressive; gene therapy; antisense therapy; livestock feed additive; cancer; autoimmune disease; transplantation;

PCR primer; ss.

Homo sapiens

11-JAN-2001.

WO200102420-A1

30-JUN-2000; 2000WO-US018269

01-JUL-1999;

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RESULT 3416
ABA94871/c
ID ABA9487
XX ABA9487
XX ABA9487
XX OB-MAY-
XX Tumour
KW Tumour
KW LUCA-2;
KW CYCOsta
XX Homo sa
X
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is given in a specification relating to an isolated and purified nucleic acid encoding a biologically active methionine adenosyltransferase II (MAT II) beta subunit polypeptide capable of modulating MAT II biological activity. The MATT II beta subunit nucleic acid and polypeptide are useful in screening assays for identifying compounds that affect or modulate MATT II biological activity. The MATT II beta subunit polypeptides have utility as feed additives for livestock. The MATT II beta subunit polypeptide, anti-MATT II beta subunit antibody and antisense oligonucleotide are useful for treating a disorder associated with MAT II biological activity. e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumour suppressor gene; chromosome 3p21.3; Gene 26; PL6; Beta*; L'LUCA-2; 123F2; Fusl; 101F6; Gene 21; SEM A3; cancer; tumour; BLU; cytostatic; gene therapy; protein therapy; RASFFIA; PCR primer; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid encoding a biologically active methionine adenosyltransferase II (MAT II) beta subunit, useful in for identifying MATT II modulating compounds which can b cancer and autoimmune diseases.
                           New tumor suppressor genes and proteins, useful for detecting, and treating various human cancers, e.g. spleen cancer, kidney lymph node cancer, small intestine cancer, blood cell cancer or pancreatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unmethylated RASSF1A promoter fragment detecting reverse
                                                                                                                                                                  WPI; 2002-179699/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABA94871 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                       10-JUL-2000; 2000US-0217112P
                                                                                                                                                                                                                                                                                                                                                                                                          10-JUL-2001; 2001WO-US021781.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200204511-A2
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17; Conserv
                                                                                                                                                                                                                 Minna J,
                                                                                                                                                                                                                                                                      UNIV TEXAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                         HEALTH
                                                                                                                                                                                                                    Roth J,
                                                                                                                                                                                                                                                                                               SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.2%;
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Pred. No. 2.4e
0; Mismatches
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                                                                                                                                                                                                                                                                            SERVICES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene 26; PL6; Beta*; LUCA-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              n screening assays be used to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                          diagnosing
                                                                                   cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human;
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ABL92904
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
07-AUG-2000;
07-AUG-2000;
11-AUG-2000;
11-AUG-2000;
05-JAN-2001;
05-JAN-2001;
08-JAN-2001;
10-JAN-2001;
10-JAN-2001;
22-JAN-2001;
23-JAN-2001;
23-JAN-2001;
09-FEB-2001;
09-FEB-2001;
15-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to tumour suppressor genes at chromosome 3p21.3 The invention relates to tumour suppressor genes at chromosome 3p21.3 (Gene 26, PL6, Beta*, LUCA-1, LUCA-2, 123F2, Fus1, 101F6, Gene 21, or SEM A3). The tumour suppressor genes play a major role in the pathogenesis of human lung cancer and other cancers. The polypeptide and polynucleotides are useful for detecting, diagnosing and treating various human cancers. These are particularly useful for inhibiting human cancers.
                                                                                                                                                                                                                                                                                                                                                                                                  Human; G protein-coupled receptor; antidiabetic; anorectic; cytostatic; immunomodulator; neuroprotective; notropic; antiparkinsonian; metabolic immunosuppressive; ophthalmological; antibacterial; virucide; fungicide; protozoacide; hypertensive; hypotensive; analgesic; osteopathic; antilizer; antiatethmatic; antiallergic; anti-HIV; antilipaemic; vaccine; antiinfertility; antiinflammatory; haemostatic; cell signal processing; cardiomyopathy; atherosclerosis; metabolic pathway modulation; cancer; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumourigenicity, suppressing tumour or restricting metastatic processes in various tumours such as lung, colon, breast, stomach, cervix, and head and neck, prostate or pancreas. In particular, the cancer is brain cancer, lung cancer, letter cancer, spleen cancer, kidney cancer, lymph node cancer, small intestine cancer, blood cell cancer, pancreatic cancer, colon cancer, stomach cancer, cervix cancer, breast cancer, endometrial cancer, stomach cancer, testicle cancer, ovarian cancer, skin cancer, head and neck cancer, esophageal cancer, ovarian cancer or bone marrow cancer. The present sequence represents a primer for detecting an unmethylated RASSFIA promoter fragment by methylation-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G protein-coupled receptor GPCR23 PCR primer SEQ ID NO:386.
                                                                                                                                                                                                                                                                      07-AUG-2001;
                                                                                                                                                                                                                                                                                                     14-FEB-2002.
                                                                                                                                                                                                                                                                                                                                      WO200212343-A2
                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL92904 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCB.
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                 2000US-0223138P.
2000US-0224613P.
2000US-0224615P.
2001US-0260073P.
2001US-0260073P.
2001US-0260073P.
2001US-0260450P.
2001US-0261156P.
2001US-0261156P.
2001US-0263338P.
2001US-0263434P.
2001US-0263434P.
2001US-0263704P.
2001US-026964P.
2001US-026964P.
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                                                                                                                                                                                                                                                                      2001WO-US024787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.4e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          metabolic,
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RESULT 3418
ABL92913
ID ABL9291
XX ABL9291
XX D6-JUN-
XX D6-JUN-
XX Human;
KW Human;
KW immunom
KW immunom
KW immunom
KW immunom
KW antiinf
KW antiinf
KW antiinf
KW cardiom
KW gene th
XX Homo sa
XX WO20021
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC neuroprotective; notropic; antiparkinsonian; analyses of operative; contropic; antiparkinsonian; analyses of operative; contropic; antiparkinsonian; anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes novel human G protein-coupled receptors (GPCR) designated GPCR1-36 from the present invention. The GPCRs can have activities such as: antidiabetic; anorectic; immunomodulator; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spytek KA,
Casman SJ,
     WO200212343-A2
                                                                                                  immunosuppressive; ophthalmological; antibacterial; virucide; fungicide; protozoacide; hypertensive; hypotensive; analgesic; osteopathic; antiulcer; antiasthmatic; antiallergic; anti-HIV; antillpaemic; vaccine; antiinfertility; antiinflammatory; haemostatic; cell signal processing; cardiomyopathy; atherosclerosis; metabolic pathway modulation; cancer; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                         Human; G protein-coupled receptor; antidiabetic; anorectic; cytostatic; immunomodulator; neuroprotective; nootropic; antiparkinsonian; metabolic;
                                                                                                                                                                                                                                                                                                                                                                                           06-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 407; 492pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New G-protein coupled receptor polypeptides and nucleic acids, useful for diagnosis, prevention or treatment of hematopoletic, neurodegenerative,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-217180/27.
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                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL92913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL92913 standard;
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                                                                                                                                                                                                                                                                                                                                      protein-coupled receptor GPCR25 PCR primer SEQ ID NO:395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           l Similarity
17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGGTAGATGGGGAAGAAATAC 3661
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Vernet CAM,
Mezes PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Taupier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 C; 9
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Ballinger RA, Shenoy SG,
Grosse WM, Alsobrook JP,
RJ, Colman SD, Szekeres E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . JC, Li
Kekuda
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CC antiasthmatic; antilipaemic; antiallergic; antiinfertility; haemostatic; cc antiasthmatic; antilipaemic; antiallergic; antiinfertility; haemostatic; cc and antiinfelammatory. They can be used in gene therapy and vaccine production. The GPCR proteins can be used for treating or preventing GPCR cc -associated disorders such as cardiomyopathy, atherosclerosis, or a cc disorder related to cell signal processing and metabolic pathway cm modulation, in humans. GPCR proteins and the polynucleotides encoding cc disease, especially cancer associated with altered levels of GPCR croteins and polynucleotides, by measuring the level of protein GCC expression or the amount of nucleic acid from a mammal and comparing it composition are also useful for identifying an agent, especially cellular cc proteins are also useful for identifying an agent, especially cellular cc candidate substance interacting with an olfactory receptor or screening of a cc andidate substance interacting with an olfactory receptor polypeptide, its fragments or variants. The present sequence represents a PCR primer cc used in the isolation of a novel human GPCR in the present invention
                                                                                         Query Match
Best Local S
Matches 17
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Casman SJ,
Burgess CE,
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22-JAN-2001;
23-JAN-2001;
01-FEB-2001;
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05-JAN-2001;
08-JAN-2001;
09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes novel human G protein-coupled receptors (GPCR) designated GPCR1-36 from the present invention. The GPCRs can have activities such as: antidiabetic; anorectic; immunomodulator; cytostatic; neuroprotective; nootropic; antiparkinsonian; analgesic; osteopathic; immunosuppressive; metabolic; ophthalmological; antibacterial; virucide; fungicide; protozoacide; hypertensive: hypertensive:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-AUG-2000;
07-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New G-protein coupled receptor polypeptides and nucleic acids, useful for diagnosis, prevention or treatment of hematopoietic, neurodegenerative,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-AUG-2000;
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                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-217180/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-AUG-2000;
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1 TGCCTCCCTTTCTGACTAAGT 21
                                                                                           l Similarity
17; Conserv
                                               TGCCTGCCTGTCTGCCTATGT 5821
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            signal transduction pathway disorders.
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2001US-0260072P
2001US-0260283P
2001US-0260450P
2001US-0261156P
2001US-0263134P
2001US-0263704P
2001US-0265704P
2001US-0265704P
2001US-0276406P
2001US-0276406P
2001US-0276406P
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Mezes PS,
                                                                                                                                                                                       B₽;
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                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001WO-US024787
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                                                                                                               0.2%;
81.0%;
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                                                                                                                                                                                       C; 3
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Ballinger RA, Shenoy SG
Grosse WM, Alsobrook JP,
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                                                                                                               Score 14.6;
Pred. No. 2
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                                                                                                                                                                                          9
                                                                                              Mismatches
                                                                                                                                                                                          Η.
                                                                                                                                                                                       0 U;
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                                                                                                                                                                                            0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Szekeres
                                                                                                                  4e+03;
                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SG,
                                                                                                                                           Length
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Kekuda R;
Gorman L;
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RESULT 3419 ABK52319/c

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RESULT 3420
ABS58924/c
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                                                                                                                Human; G-protein coupled receptor; GPCR; cardiomyopathy; atherosclerosis; diabetes; cell signal processing; metabolic pathway modulation; cancer; adenocarcinoma; lymphoma; prostate cancer; uterus cancer; asthma; immune response; neurodegenerative disorder; inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     smooth muscle cell proliferation and a method for screening a substance having inhibitory activity on vascular smooth muscle cell proliferation. This sequence represents a PCR primer associated with the method of measuring, and screening for compounds responsible for, inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Measurement of the inhibitory activity for vascular smooth proliferation, and a method for screening a substance with activity for vascular smooth muscle cell proliferation.
                                                                                                       immune response; neurodege Crohn's disease; multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vascular smooth muscle; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vascular smooth muscle cell proliferation associated PCR primer #7.
                       01-AUG-2002
                                               WO200259313-A2
                                                                      Homo sapiens.
                                                                                                                                                                                Human G-protein coupled receptor, reverse primer
                                                                                                                                                                                                          05-NOV-2002
                                                                                                                                                                                                                                                          ABS58924
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; Page 22; 24pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-AUG-2000; 2000JP-00235459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-SEP-2000; 2000JP-00284973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP2002112798-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK52319
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                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                            4646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002-448760/48
                                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                            TGGAATTTCCTCTTTGAGGAG 4666
                                                                                                                                                                                                                                                                                                                                                                                                                  22 BP; 8 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                     TGGATTTGCCTCATTGTGGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                             smooth
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    describes a measurement of inhibitory activity on
                                                                                                                                                                                                                                                                                                                                                                                                                                           muscle cell proliferation
                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                               81.0%;
                                                                                                                                                                                                                                                                                                                                                                                             0.2%;
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                                                                                                                                                                                                                                                         22
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                                                                                                       sclerosis; Albright hereditary osteodystrophy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                Score 14.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                               .4e+03
                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proliferation inhibitor;
                                                                                                                                                                                                                                                                                                                                                                    4.
                                                                                                                                                                                #24.
                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                             22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  muscle cell
inhibitory
                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         on vascular
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18-DEC-2001; 2001WO-US049394.

BXAXAXB

AAD43245 AAD43245

standard;

DNA;

22

ВP

0

14-NOV-2002

(first entry

Antisense oligonucleotide R51AS6.

RESULT 3421

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                                                                                                                               CC polypeptides and polynucleotides. The GPCR polypeptide, GPCR nucleic acid CC and antibody are useful for treating, preventing or alleviating a GPCR-CC associated disorder or a pathological state in a subject, particularly a CC human. In particular, the disorder is cardiomyopathy, atherosclerosis, CC diabetes, or a disorder related to cell signal processing and metabolic comparts of the presence of or predisposition to a disease associated cC pathway modulation. The GPCR polypeptide and nucleic acid are also useful cC for diagnosing the presence of or predisposition to a disease associated cC with altered levels of GPCR, particularly cancer. The GPCR nucleic acid capilications for disorders associated with aberrant GPCR expression or activity. The DNA encoding the protein is useful in gene therapy for CC treating the above conditions. Furthermore, the nucleic acids and CC polypeptides are useful in treating adenocarcinoma. lymphoma, prostate cancer, uterus cancer, immune response, neurodegenerative disorders, Cabrima, inflammatory disorders, Crohn's disease, multiple sclerosis or CC astright hereditary osteodystrophy. These are also useful in developing a powerful assay system for functional analysis of various human disorders, CBCPB_Coding_series and arches of the investion thuman disorders.
                                                                               Query Match
Best Local
                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JAN-2001;
12-JAN-2001;
24-JAN-2001;
08-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-DEC-2000;
21-DEC-2000;
04-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Casman S
Gerlach
                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in gene therapy, atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-FEB-2001;
14-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peyman JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-APR-2001;
19-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CURA-)
                    3921 CTCTTGGCTTCTTTTCTCCCT 3941
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein coupled receptor polypeptides and polynucleotides, useful ne therapy, particularly for treating or preventing cardiomyopathy, osclerosis, diabetes, multiple sclerosis, Crohn's disease or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ballinger RA, Pao
n SJ, Vernet CAM,
n S, Smithson G,
n JA, Ellerman K,
21
                                                                17;
                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page
                                                                                                                               22
CTCTTTGGTTCTTTGATCCCT 1
                                                                                                                             BP; 12 A;
                                                                                                                                                            sequences, primers and probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; 2000US-0258635P.
; 2001US-0259743P.
; 2001US-0259741P.
; 2001US-0260718P.
; 2001US-0263689P.
; 2001US-0267464P.
; 2001US-02779464P.
; 2001US-0277946P.
; 2001US-0277945P.
; 2001US-0285718P.
; 2001US-0285718P.
; 2001US-0285718P.
; 2001US-0285718P.
; 2001US-029327P.
; 2001US-0312902P.
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          relates to novel isolated G-protein coupled receptor (GPCR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        300; 685pp;
                                                                          0.2%;
                                                                                                                           3 C; 6 G; 1 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Padigaru M, Kekuda R,
M, Shenoy SG, Gusev V,
G, Stone DJ, Sciore P,
K, Gangolli EA, Millet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English
                                                             0,
                                                                              Score 14.6;
Pred. No. 2
                                                              Mismatches
                                                                            .4e+03
                                                                                             DB 1;
                                                                                                                                                               o
H
                                                                                                                               0 Other;
                                                                                                                                                               the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Colman SD, Spyt
Malyankar UM,
Macdougall JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          н
                                                                                           Length
                                                              Indels
                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spytek KA;
                                                             0;
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Gunther E;
                                                             Gaps
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ABL35690
ID ABL3569
XX ABL3569
XX ABL3569
XX ABL3569
XX Inmunos
XX DNA/RNA
KW Infecti
KW inmunos
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                                                                                                                                                                                                                                                                                                                           Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a method for inhibiting or reducing tumour cell proliferation in an individual in vivo. The method comprising contacting a tumour cell in vivo with a Rad51 inhibitor and a polynucleotide capable of expressing functional p53 protein. The method is useful for inhibiting or reducing tumour cell proliferation in an individual in vivo. The method is useful for treating hyperproliferative disorders, especially cancer (such as Hodgkin's disease, squamous cell carcinoma and leukaemia), premature aging, autoimmune disease, arthritis, graft rejection, inflammatory bowel disease, and proliferation induced after medical procedures such as surgery and angioplasty. The invention is useful in gene therapy. The present sequence is an antisense oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumour cell proliferation; Rad51 inhibitor; p53 protein; premature aging; hyperproliferative disorder; Hodgkin's disease; squamous cell carcinoma; leukaemia; autoimmune disease; cancer; graft rejection; angioplasty; inflammatory bowel disease; immunosuppressive; gene therapy; arthritis; antisense; phosphorothicate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inhibiting/reducing tumor cell proliferation in individual in vivo, for treating cancer, arthritis, involves contacting tumor cell in vivo with Rad51 inhibitor, and polynucleotide expressing functional p53 protein.
                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 5; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-635686/68
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                                          DNA/RNA hybrid;
                             infection; allergy;
                                                                          Immunostimulatory oligonucleotide SEQ ID NO: 616.
                                                                                                                                                                       ABL35690 standard;
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                                                                                                                                                                                                                                                                                 3477
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                                                                                                                                                                                                                                                                                                                l Similarity
17; Conser
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                                                                                                                                                                                                                                                                               CCCTAGTAATACTTAAGGCAC 3497
                                                                                                                                                                                                                                                                                                                                                                            22
                                                                                                                                                                                                                                                    CCCAAGTCATTCCTAAGGCAC 21
                                                                                                                                                                                                                                                                                                                                                                           BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reddy G;
                                                                                                         (first entry)
                                          phosphorothicate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
             gy; cancer; hy, antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                            Α,
                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                           0.2%;
                                                                                                                                                                                                                                                                                                                                                                            9 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            _base= OTHER
                                                                                                                                                                       22
                                                                                                                                                                       ВÞ
hypersensitivity; bio-warfare; ic; cytostatic; antimicrohic;
                                                                                                                                                                                                                                                                                                                                                                         G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                              Score 14.6; E
Pred. No. 2.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                               2.4e+03;
                                                                                                                                                                                                                                                                                                                                            DB 1; Length 22;
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SXXXXXXXXXXXX

FGF10 amplifying reverse RT

PCR primer.

10-SEP-2002 AAD38021;

(first entry)

Lung cancer; homeodomain; homeobox; HOX; wingless/int-1; WNT; tumour; diagnosis; prognosis; therapeutic; real time PCR; RT-PCR; FGF10; prim

FGF10; primer;

Unidentified

밁 S

AAD38021

AAD38021 standard;

DNA;

22 ВP

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Matches
                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                   The present invention relates to an immunostimulatory composition, which comprises at least one oligonucleotide comprising both an RNA region and a DNA region. The composition is useful for enhancing an immune response or inducing cytokines. It can be used as a vaccine adjuvant and in treating diseases, including pathogenic infection, (non-)malignant tumours (e.g. cancers of the brain, lung, ovary, breast, prostate or colon, or carcinomas and sarcomas), autoimmune diseases or allergies (e.g. allergic rhinitis, hay fever or food allergies). Lyme disease, preventing or ameliorating, the symptoms resulting from exposure to a biowarfare agent, e.g. Ebola, Anthrax or Listeria. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New immunostimulatory compositions comprising RNA/DNA hybrid oligonuclectides, useful for enhancing an immune response or cytokines, particularly for treating diseases, e.g. cancer, s HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunosuppressive; protozoacide; virucide; hepatotropic; gene therapy;
antiinflammatory; antibacterial; ss.
                                                                                                                                                                                                             Sequence 22 BP; 0 A; 3 C; 2 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                     the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 11; Page 63; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mond JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-2001; 2001WO-US018276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOS-)
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                            4463 CTTTTTTTTTTTTTTTTTT 4483
                                                                                                        l Similarity
17; Conserv
N
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CGTTTGTTCTCTTTTTTTTT
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                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0209797P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "optionally thymidine is replaced by uracil to form RNA or DNA/RNA hybrids. Thymidine is linked to at least one other base through a ribose sugar"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Klinman
                                                                                                                                 81.0%;
                                                                                                                                                             0.2%;
                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DW.
                                                                                                                                    Score 14.6;
Pred. No. 2
                                                                                                          Mismatches
                                                                                                                                      .4e+03
                                                                                                                                                             DB 1; Length 22;
                                                                                                          Indels
                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       allergy
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                                                                                                          Gaps
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RESULT 3424
ABS78424
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to evaluating a lung cell sample of a lung cancer patient comprising screening for expression of a homeodomain containing homeobox (HOX) or wingless/int-1 (MVT) gene and comparing the screen to a sample of non-malignant lung cell. The method is useful for evaluating a lung cell sample of a lung cancer patient. The method is useful for determining the prognosis of a lung cancer and for assessing a lung cancer for a tendency to metastasise, where decreased expression of a WNT gene or HOXAl gene, or increased expression of HOX gene other than HOXAl, or both, indicates poor prognosis, increased rate of tumour growth, and increased tendency to metastasise. The method is useful for obtaining data for diagnosis, to determine clinical prognosis, including rate of growth, tendency to metastasise to assess the efficacy of treatment and to determine the efficacy of a therapeutic agent. The present sequence is a real time (RT)-PCR primer used to amplify FGF10. This sequence is used in the expendition of the author of the present sequence is a real time (RT)-PCR primer used to amplify FGF10.
                                                                                                                                                                                   Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; soriasis; tumour metastasis; precancerous lesion; rheumatoid arthritis; soriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Evaluating a lung cell sample of a lung cancer patient for diagnosis to determine clinical prognosis including tendency to metastasize comprises screening for expression of homeodomain containing genes HOX or WNT.
                                                                                                                                                                                                                                                                                                                 13-DEC-2002
                                                 WO200253141-A2
                                                                                  Synthetic
                                                                                                                                   angiofibroma; wound granulation; intestinal
                                                                                                                                                    rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac j
                                                                                                                                                                                                                                                                           Angiogenesis inhibitory oligonucleotide #908
                                                                                                                                                                                                                                                                                                                                                                                     ABS78424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-OCT-2000; 2000US-0239596P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-OCT-2001; 2001WO-US031960
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCCTTTTCTTCTATTCACTC 5760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TCCATTTTCCTCTATCCTCTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37pp;
                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.6;
Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4;
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                                                                                                                                     atherosclerosis;
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                                                                                                                                                       joint;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retnopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubcosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque meovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                  (CSIR )
(DAIR-)
(AGRI-)
                                                                                                                                                                                                                                                 Virus-encoded coat protein; primer; AMV; CYVV; WCMV; leguminous plant; viral replicase; intron ribonucleic acid; iRNA; hairpin RNA; virucide; pathogenic plant virus; SSU promoter; 35S promoter; gene therapy; RT-PCR;
                                                                                                                  16-NOV-2001; 2001WO-AU001496.
                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                             Alfalfa mosaic
                                                                                                                                                                                                                                                                                                        Alfalfa mosaic virus DNA RT-PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                       ABK97133 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; Page 35; 276pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inhibiting angiogenesis in a subject, antiangiogenic nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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                                                                                       17-NOV-2000; 2000AU-00001558
                                                                                                                                             23-MAY-2002.
                                                                                                                                                                       WO200239808-A1
                                                                                                                                                                                                                                                                                                                                   07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                              АВК97133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-DEC-2001; 2001WO-US048458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5328
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                                    COMMONWEALTH SCI & IND RES ORG. DAIRY RES & DEV CORP. AGRIC VICTORIA SERVICES PTY LTI
                                                                                                                                                                                                                                      transcriptase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCTCTTTGCCTCACTCTCTC 5348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                  RES & DEV
                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                              Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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Pred. No. 2
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Chu PWG,

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ABN84449
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Best Local Similarity
Matches 17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dysfunctional viral replicase or an intron ribonucleic acid (IRNA) comprising a hairpin RNA to the plant so the plant is immune to the plant virus under field conditions. The invention also relates to a method for transforming a leguminous plant by introducing to a leguminous plant cell, tissue or organ, an isolated nucleic acid molecule comprising an SU promoter or a 350 promoter operably linked to a nucleic acid encoding a virus-encoded coat protein and regenerating a transformed leguminous plant from the plant cell, tissue or organ, where the transformed leguminous plant is immune to a pathogenic plant virus. The method is useful for producing a leguminous plant with enhanced viral resistance by crossing two parent plants each having enhanced viral resistance or immunity against one or more different viruses. This sequence represents a reverse transcriptase PCR (RT-PCR) primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conferring immunity to pathogenic plant virus on leguminous plant by introducing nucleic acid sequence encoding virus-encoded coat protein, dysfunctional viral replicase or iRNA comprising hairpin RNA, to plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bcr/abl forward PCR primer used in whole cell RT-PCR.
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                                                                               Reducing the activity of a ribonuclease, useful for inhibiting tumor growth or for removing RNA binding proteins from a solution, comprises mixing a sample containing an RNase with a preparation containing at
                                                                                                                                                                                                                                                                                                                                                                                                         28-NOV-2000;
28-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA polymer; ribonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-OCT-2002
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                                                                                                                                                                                                                WPI; 2002-599806/64.
                                                                                                                                                                                                                                                                            Shultz JW, Lewis MK,
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                                                                                                                                                                                                                                                                                                                                          (PROM-) PROMEGA CORP
                                                    least one RNA polymer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2238 CCAGATACTCCATTATGAGCT 2258
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2001US-00995912
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Example 7; Page 45; 61pp; English

WPI; 2002-643487/69

Casman SJ,

Edinger SR,

Ellerman K,

Smithson

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Kekuda

CURAGEN

CORP.

Muralidhara

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The present sequence is that of a bcr/abl forward PCR primer. The primer CC was used in an example from the invention demonstrating the inhibition of CC RNase enzymes in RT-PCR reactions using polyg or polyI attached to CC resins. RT-PCR was conducted without prior RNA isolation using K562 CC (human erythroleukaemia cell line) whole cell lysates. The reverse primer CC for the PCR is given in ABNG4450, Results were compared with the effect CC of adding RNASIN ribonuclease inhibitor. Addition of RNASIN to cells CC during lysis allowed for the sensitive detection of the bcr/abl signal CC down to as low as I cell, with increasing signal intensity with CC increasing cell number. The addition of 1 ul, but not 3 ul, of polyG CC cells (with or without spin). The signal in the presence of polyG wase than with RNASIN. It is concluded that RNA polymers can replace CC RNASIN in single-tube, whole cell RT-PCR. The example illustrates the CC present invention, which relates to compositions and methods for using CC RNA polymers to inhibit RNase enzymes (e.g. those involved in CC angiogenesis such as in tumour growth and proliferation, and in other CC involved in disease states

CC involved in disease states
Sequence 22 BP; 6 A; 6 C; 7 G; 3 T; 0 U;
  0 Other;
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RESULT 3427
ABS64779
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Best Local :
                                                                                                                                                                                                                                           G protein coupled receptor related protein; 88; primer; human; GPCR; cardiomyopathy; atherosclerosis; diabetes; cancer; stroke; PCR; Von Hippel-Lindau syndrome; Alzheimer's disease; tuberous sclerosis; hypercalcaemia; Parkinson's disease; Huntington's disease; cerebral paley; epilepsy; Lesch-Nyhan syndrome; multiple sclerosis; ataxia-telangiectasia; leukodystrophy; addiction; anxiety; depression; pain; obesity; Crohn's disease; osteoporosis; haemophilia; asthma; inflammatory bowel disease; infertility; hypertension; scleroderma; arthritis; human immunodeficiency virus; autoimmune disease; HIV; infection; graft-versus-host disease.
                                                                                     03-JAN-2001; 2001US-0259552P
09-JAN-2001; 2001US-0260544P
20-MAR-2001; 2001US-0277405P
                                                                                                                                         03-JAN-2002; 2002WO-US000056.
                                                                                                                                                                                                                                                                                                                                                                                                GPCR15 real time forward PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS64779;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABS64779 standard; DNA; 22
                                                                                                                                                                    22-AUG-2002.
                                                                                                                                                                                               WO200264793-A2
                                                                                                                                                                                                                         Homo sapiens.
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17; Conserv
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